

GenCore version 5.1.8  
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OM protein - protein search, using sw model

Run on: May 12, 2006, 15:15:11 ; Search time 226 Seconds  
(without alignments)  
115.507 Million cell updates/sec

Title: US-08-870-762b-1  
Perfect score: 202  
Sequence: 1 KCNTATCATQRLANFLVHSSNNFGPILPTVNGSNTY 37

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 2166443 seqs, 705528306 residues

Total number of hits satisfying chosen parameters: 2166443

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 100 summaries

Database : UniProt 05.80:\*  
1: uniprot\_sprot:\*  
2: uniprot\_crembl:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	187	92.6	93	1 IAPP_MOUSE	P12968 mus musculus
2	187	92.6	93	1 IAPP_RAT	P12960 rattus norv
3	184	91.1	37	1 IAPP_CRICR	P19690 cricetus
4	184	91.1	92	1 IAPP_MESAU	P23442 mesocricetu
5	178	88.1	89	1 IAPP_HUMAN	P10997 homo sapien
6	171	84.7	89	1 IAPP_FELCA	P12667 felis silve
7	169	83.7	89	1 IAPP_CANFA	P17116 canis fami
8	160	79.2	91	1 IAPP_OCTDR	P22889 octodon deg
9	153	75.7	135	2 Q90743_CHICK	Q90743 gallus gall
10	151	74.8	92	1 IAPP_CAVPO	P12966 cavia porce
11	148	73.3	67	1 IAPP_RABIT	Q07334 oryctolagus
12	143	70.8	51	2 Q4TB97_TETNG	Q4TB97 tetraodon n
13	132	65.3	66	2 Q9BEP0_ERIEU	Q9BEP0 erinaceus e
14	120	59.4	32	1 IAPP_SAGOE	Q28934 saginus ce
15	107	53.0	32	1 IAPP_PIG	Q29119 sus scrofa
16	104	51.5	32	1 IAPP_SHEEP	Q28605 ovis aries
17	96	47.5	23	1 IAPP_LABEU	Q28733 lepus europ
18	96	47.5	37	1 IAPP_BOVIN	P31888 rana ridibu
19	96	47.5	32	1 CALCR_RANRI	P10286 gallus gall
20	96	47.5	125	1 CALCA_CHICK	P81564 phyllomedus
21	94	46.5	115	1 CALCR_PHYRI	Q6d9j brachydanio
22	92	45.5	126	2 Q6DGU5_BRARE	Q6d9j brachydanio
23	91	45.0	25	2 Q9BEE1_MACRO	Q9BEE1 macropus ru
24	90	44.6	52	2 P79814_ONCGO	P79814 oncorhynch
25	90	44.6	126	2 Q8QFT9_FUGRU	Q8QFT9 fuqua rubrip
26	90	44.6	184	2 Q4S167_TETNG	Q4S167 tetraodon n
27	89	44.1	56	2 Q92164_ONCSP	Q92164 oncorhynch
28	89	44.1	128	1 CALCA_CANFA	Q9myv1 canis fami
29	88	43.6	51	2 Q4S173_TETNG	Q4S173 tetraodon n
30	86	42.6	127	1 CALCB_HUMAN	P10092 homo sapien
31	86	42.6	127	2 Q56910_HUMAN	Q56910 homo sapien

32	85	42.1	37	1 CALCA_SHEEP	P30981 ovis aries
33	85	42.1	44	2 Q4THN5_TETNG	Q4thn5 tetraodon n
34	85	42.1	50	2 Q66VC1_RAT	Q66vc1 rattus norv
35	85	42.1	128	1 CALCA_MOUSE	Q991a0 mus musculu
36	85	42.1	128	1 CALCA_RAT	P01256 rattus norv
37	84	41.6	52	2 Q8WNX3_CALJA	Q8wnx3 callithrix
38	84	41.6	53	2 Q90YC3_PAROL	Q90yc3 paralithrix
39	84	41.6	128	1 CALCA_HUMAN	P06881 homo sapien
40	84	41.6	134	1 CALCB_RAT	P10093 rattus norv
41	82	40.6	130	1 CALCB_MOUSE	P30880 mus musculu
42	81	40.1	37	1 CALCA_PIG	Q8wnx2 callithrix
43	81	40.1	53	2 Q8WNX2_CALJA	Q8wnx2 equus cabal
44	80	39.6	129	1 CALCB_HORSE	Q9n0c3 equus cabal
45	79	39.1	60	2 Q9GLK1_RABIT	Q9glk1 oryctolagus
46	79	39.1	67	2 Q9GLK2_RABIT	Q9glk2 oryctolagus
47	74	36.6	125	2 Q75V95_BOVIN	Q75v95 bos taurus
48	71	35.1	126	2 Q766Y6_PIG	Q766y6 sus scrofa
49	70	34.7	125	2 Q862B1_PIG	Q862b1 sus scrofa
50	67	33.2	117	2 Q766Y7_PIG	Q766y7 sus scrofa
51	66	32.7	127	1 CALCA_HORSE	Q9n0c2 equus cabal
52	61.5	30.4	178	1 Q9UAV9_CAEEL	Q9uav9 caenorhabd
53	60	29.7	590	2 Q7PX05_ANGOA	Q7px05 anopheles g
54	59.5	29.5	179	2 Q60WM3_CAEER	Q60wm3 caenorhabd
55	58	28.7	338	2 Q9LSG7_ARATH	Q9lsq7 arabidopsis
56	57	28.2	248	2 Q812V2_PLA7	Q812v2 plasmodium
57	56.5	28.0	430	2 Q9SZA6_ARATH	Q9s2a6 arabidopsis
58	56.5	28.0	559	2 Q0Z263_CAEEL	Q0z263 caenorhabd
59	56.5	28.0	962	2 Q9LRX1_ARATH	Q9lrg7 arabidopsis
60	56	27.7	127	2 Q75V94_CANFA	Q75v94 canis fami
61	56	27.7	149	2 Q5W957_GILTI	Q5w957 dioscorea p
62	56	27.7	8402	2 QARE89_TETNG	Q4r89 tetraodon n
63	55.5	27.5	842	2 Q7R6K2_GIALA	Q7r6k2 giardia lam
64	55	27.2	347	2 Q91XF4_MOUSE	Q91xf4 mus musculu
65	55	27.2	549	2 Q6ZSA6_HUMAN	Q6zsa6 homo sapien
66	55	27.2	866	2 Q7PP39_RAT	Q7cp39 rattus norv
67	55	27.2	1050	1 TIFP1A_HUMAN	O1564 homo sapien
68	55	27.2	1051	1 TIFP1A_MOUSE	O64127 mus musculu
69	54.5	27.0	140	1 Y14K_CSNV	P18920 chloris str
70	54	26.7	300	2 Q6FAD1_ACTIAD	Q6fad1 actinobact
71	54	26.7	354	2 Q4TQ45_9SPHN	Q4tq45 erythrobact
72	54	26.7	444	2 Q7ABE2_ECO57	Q7abe2 escherichia
73	54	26.7	444	2 Q8X922_ECO57	Q8x922 escherichia
74	54	26.7	917	1 N1A1_ARATH	P11832 arabidopsis
75	54	26.7	1242	2 Q6CNK0_KIULA	Q6cnk0 kluyveromyc
76	53.5	26.5	219	2 Q7ZWS3_LBPIK	Q7zws3 leptospira
77	53.5	26.5	428	2 Q70VZ1_CIOIN	Q70vz1 ciona intes
78	53.5	26.5	659	2 Q4H3K2_CIOIN	Q4h3k2 ciona intes
79	53.5	26.5	761	2 Q4RN84_TETNG	Q4rtn8 tetraodon n
80	53	26.2	217	2 Q6RTJ9_HUMAN	Q6rtj9 homo sapien
81	53	26.2	346	2 Q4IDQ4_GIBZE	Q4idq4 gibberella
82	53	26.2	392	2 Q9L104_ORYSA	Q9l104 oryza sativ
83	53	26.2	509	2 Q5SN48_CRYNBE	Q5sn48 cryptococcu
84	53	26.2	549	2 Q5SCZ7_DICDI	Q5scz7 dictyosteli
85	53	26.2	1093	2 Q54GV1_DICDI	Q54gv1 dictyosteli
86	53	26.2	1339	2 Q5CNM0_CRYHO	Q5cnm0 cryptospori
87	53	26.2	2653	2 Q8IHX2_PLA7	Q8ihx2 plasmodium
88	52.5	26.0	233	2 Q9YER4_ARPE	Q9yer4 aeropyrum p
89	52.5	26.0	341	2 Q4X2E1_PLACH	Q4x2e1 plasmodium
90	52	25.7	407	2 Q8LSK1_MAIZE	Q8lsk1 zea mays (m
91	52	25.7	149	2 Q5W956_GILTI	Q5w956 dioscorea p
92	52	25.7	310	2 Q733U2_BACCI	Q733u2 bacillus ce
93	52	25.7	310	2 Q81YB3_BACAN	Q81yeb3 bacillus an
94	52	25.7	315	2 Q8HFP4_BACHK	Q8hfp4 bacillus th
95	52	25.7	323	2 Q9BKX8_CAEEL	Q9bkx8 caenorhabd
96	52	25.7	347	2 Q4QM34_HAE18	Q4qm34 haemophilus
97	52	25.7	424	1 SYS_PROMP	Q700r7 pneumocloro
98	52	25.7	434	2 Q6HVN6_BACAN	Q6hvn6 bacillus an
99	52	25.7	434	2 Q637S1_BACAZ	Q637s1 bacillus ce
100	52	25.7	499	2 Q4XV12_PLACH	Q4xv12 plasmodium

## ALIGNMENTS

```

RESULT 1
IAPP_MOUSE
ID IAPP_MOUSE STANDARD; PRT; 93 AA.
AC P12968;
DT 01-OCT-1989 (Rel. 12, Created)
DT 01-OCT-1989 (Rel. 12, Last sequence update)
DT 13-SEP-2005 (Rel. 48, Last annotation update)
DE Islet amyloid polypeptide precursor (Diabetes-associated peptide)
DE (DAP) (Amylin).
GN Name=Iapp;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
OC Muridae; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
[1]
NP NUCLEOTIDE SEQUENCE.
RX MEDLINE=89345542; PubMed=2668946;
RA Nishi M., Chan S.J., Nagamatsu S., Bell G.I., Steiner D.F.;
RT "Conservation of the sequence of islet amyloid polypeptide in five
RT mammals is consistent with its putative role as an islet hormone.";
RL Proc. Natl. Acad. Sci. U.S.A. 86:5738-5742(1989).
[2]
NP NUCLEOTIDE SEQUENCE.
RX STRAIN=DBA/2J; TISSUE=Liver;
RX MEDLINE=97424750; PubMed=9278863; DOI=10.1677/jme.0.0190079;
RA Ekawa K., Nishi M., Ohagi S., Sanke T., Nanto K.;
RT "Cloning of mouse islet amyloid polypeptide gene and characterization
RT of its promoter.";
RL J. Mol. Endocrinol. 19:79-86(1997).
[3]
NP NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
RX STRAIN=C57BL/6J; TISSUE=Thymus;
RX MEDLINE=22286257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Klausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shemen G.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marziani K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ueda T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loggellano N.A., Peters G.J., Abramson R.D., Mullaby S.J.,
RA Boesik S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunnarsson P.H.,
RA Richardson S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahy J., Helton E., Kettelman M., Madan A., Rodriguez S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Butlerfield Y.S.N., Krzywinski M.I., Skalska U., Smalhus D.E.,
RA Scherch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
[4]
NP PROTEIN SEQUENCE OF 38-74.
RX MEDLINE=89325677; PubMed=2666169; DOI=10.1016/0014-5793(89)81467-X;
RA Bertholtz C., Christmannson L., Engstroem U., Rorsman F., Svensson V.,
RA Johnson K.H., Westermark P.;
RT "Sequence divergence in a specific region of islet amyloid polypeptide
RT (IAPP) explains differences in islet amyloid formation between
RT species.";
RL FEBS Lett. 251:261-264(1989).
[5]
NP FUNCTION: Selectively inhibits insulin-stimulated glucose
NP utilization and glycogen deposition in muscle, while not affecting
NP adipocyte glucose metabolism.
[6]
NP SUBCELLULAR LOCATION: Secreted.
[7]
NP SIMILARITY: Belongs to the calcitonin family.
[8]
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CC removed.
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DR EMBL; M25389; AAA37874.1; -; mRNA.
DR EMBL; D31820; BAA22051.1; -; Genomic DNA.
DR EMBL; BC027527; AAH27527.1; -; mRNA.
DR PIR; C3542; C3542.
DR Ensembl; ENSMUSG0000041681; Mus musculus.
DR MGI; MGI:96382; IAPP. Mus musculus.
DR GO; GO:0005615; C:extracellular space; TAS.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin B.
DR Pfam; PF00214; Calc CGRP IAPP; I.
DR PRINTS; PR00817; CALCITONINB.
DR PRINTS; PR00818; ISLETAMYLOID.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
KW Amidation; Amyloid; Cleavage on pair of basic residues;
KW Direct protein sequencing; Hormone; Signal.
FT SIGNAL 1 23 Potential.
FT PROPEP 24 35 Islet amyloid polypeptide.
FT PEPTIDE 38 74
FT PROPEP 78 93
FT MOD_RES 74 74
FT DISULFD 39 44 Tyrosine amide (G-75 provides amide
FT SEQUENCE 93 AA; 10022 MW; B15DBBC81475B15 CRC64;
                    group).
Query Match 92.6%; Score 187; DB 1; Length 93;
Best local Similarity 91.9%; Pred. No. 2,2e-18;
Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
OY 1 KNTATCATGTPLANELVHSSNNRGPILPPTVNSNTY 37
DB 38 KNTATCATGTPLANELVHSSNNRGPILPPTVNSNTY 74
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RESULT 2
IAPP_RAT
ID IAPP_RAT STANDARD; PRT; 93 AA.
AC P12969;
DT 01-OCT-1989 (Rel. 12, Created)
DT 01-OCT-1989 (Rel. 12, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Islet amyloid polypeptide precursor (Diabetes-associated peptide)
DE (DAP) (Amylin).
GN Name=Iapp;
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
OC Muridae; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
[1]
NP NUCLEOTIDE SEQUENCE.
RX MEDLINE=89345542; PubMed=2668946;
RA Nishi M., Chan S.J., Nagamatsu S., Bell G.I., Steiner D.F.;
RT "Conservation of the sequence of islet amyloid polypeptide in five
RT mammals is consistent with its putative role as an islet hormone.";
RL Proc. Natl. Acad. Sci. U.S.A. 86:5738-5742(1989).
[2]
NP NUCLEOTIDE SEQUENCE.
RX MEDLINE=89240689; PubMed=2654937;
RA Leffert J.D., Newgard C.B., Okamoto H., Milburn J.L., Luskey K.L.;
RT "Rat amylin: cloning and tissue-specific expression in pancreatic
RT islets.";
RL Proc. Natl. Acad. Sci. U.S.A. 86:3127-3130(1989).
[3]
NP NUCLEOTIDE SEQUENCE.
RX STRAIN=WAR; TISSUE=Liver;
RX MEDLINE=91027936; PubMed=2223885; DOI=10.1016/0167-4781(90)90210-S;
RA van Marfeld A.D.M., Mosselman S., Hoepfner J.W.M., Zandberg J.,
RA van Teeffelen H.A.A.M., Baas P.D., Lips C.J.M., Jansz H.S.;

```

RT "Islet amyloid polypeptide: structure and upstream sequences of the  
 RT IAPP gene in rat and man.";  
 RL Biochim. Biophys. Acta 1087:235-240(1990).  
 RN (4)  
 RP PROTEIN SEQUENCE OF 38-74.  
 RX MEDLINE=90026410; PubMed=2679555;  
 RA Asai J., Nakazato M., Kangawa K., Matsukura S., Matsuo H.;  
 RT "Isolation and sequence determination of rat islet amyloid  
 RT polypeptide.";  
 RL Biochem. Biophys. Res. Commun. 164:400-405(1989).  
 RN (5)  
 RP PROTEIN SEQUENCE OF 38-74.  
 RX MEDLINE=90290528; PubMed=2357234;  
 RA Asai J., Nakazato M., Miyazato M., Kangawa K., Matsuo H.,  
 RA Matsukura S.;  
 RT "Regional distribution and molecular forms of rat islet amyloid  
 RT polypeptide.";  
 RL Biochem. Biophys. Res. Commun. 169:788-795(1990).  
 RN [6]  
 RP NUCLEOTIDE SEQUENCE OF 38-74.  
 RX MEDLINE=89325677; PubMed=2666169; DOI=10.1016/0014-5793(89)81467-X;  
 RA Betsholtz C., Christmansson L., Engstrom U., Korsman F., Svensson V.,  
 RA Johnson K.H., Westermarck P.;  
 RT "Sequence divergence in a specific region of islet amyloid polypeptide  
 RT (IAPP) explains differences in islet amyloid formation between  
 RT species.";  
 RL FEBS Lett. 251:261-264(1989).  
 CC -1- FUNCTION: Selectively inhibits insulin-stimulated glucose  
 CC utilization and glycogen deposition in muscle, while not affecting  
 CC adipocyte glucose metabolism.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- TISSUE SPECIFICITY: Abundant in the islets of Langerhans but is  
 CC not present in the brain or seven other tissues examined.  
 CC -1- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
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 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 CC EMBL: M25390; AAA41359.1; -; mRNA.  
 CC EMBL: J04544; AAA40730.1; -; mRNA.  
 CC EMBL: X52820; CAA37003.1; -; Genomic DNA.  
 CC EMBL: X52821; CAA37003.1; JOINED; Genomic DNA.  
 CC PIR: S13566; TCRTA.  
 CC Ensembl: ENSRNOG0000012417; Rattus norvegicus.  
 DR RGD: 2654; IAPP.  
 DR InterPro: IPR000443; Amylin.  
 DR InterPro: IPR001693; Calcitonin-like.  
 DR InterPro: IPR002163; Calcitonin-B.  
 DR Pfam: PF00214; Calc\_CGRP\_IAPP; 1.  
 DR PRINTS: PR00817; CALCITONINB.  
 DR PRINTS: PR00818; ISLETAMYLID.  
 DR SMART: SM00113; CALCITONIN; 1.  
 DR PROSITE: PS00258; CALCITONIN; 1.  
 DR AmiAdacton; Amyloid; Cleavage on pair of basic residues;  
 KW Direct protein sequencing; Hormone; Signal.  
 FT SIGNAL 1 23  
 FT PROPEP 24 35  
 FT PEPTIDE 36 74  
 FT PROPEP 76 93  
 FT MOD\_RES 74 74  
 FT MOD\_RES 74 74  
 FT DISULFID 39 44  
 FT SEQUENCE 93 AA; 10015 MW; 5A76C92B624DA962 CRC64;  
 Query Match 92.6%; Score 187; DB 1; Length 93;  
 Best Local Similarity 91.9%; Pred. No. 2, 2e-18;  
 Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Db 38 KCNTATCATQRLANFLVRSNNLGPVLPPTVNGSNTY 74  
 RESULT 3  
 IAPP CRIGR STANDARD; PRT; 37 AA.  
 AC P19850;  
 DT 01-FEB-1991 (Rel. 17, Created)  
 DT 01-FEB-1991 (Rel. 17, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Islet amyloid polypeptide (Amylin).  
 GN Name=IAPP;  
 OS Cricetus griseus (Chinese hamster).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;  
 OC Muridae; Cricetidae; Cricetinae; Cricetulus.  
 OX NCBI\_Taxid=10029;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=89325677; PubMed=2666169; DOI=10.1016/0014-5793(89)81467-X;  
 RA Betsholtz C., Christmansson L., Engstrom U., Korsman F., Svensson V.,  
 RA Johnson K.H., Westermarck P.;  
 RT "Sequence divergence in a specific region of islet amyloid polypeptide  
 RT (IAPP) explains differences in islet amyloid formation between  
 RT species.";  
 RL FEBS Lett. 251:261-264(1989).  
 CC -1- FUNCTION: Selectively inhibits insulin-stimulated glucose  
 CC utilization and glycogen deposition in muscle, while not affecting  
 CC adipocyte glucose metabolism.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
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 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 CC PIR: S05037; S05037.  
 DR InterPro: IPR000443; Amylin.  
 DR InterPro: IPR001693; Calcitonin-like.  
 DR InterPro: IPR002163; Calcitonin-B.  
 DR Pfam: PF00214; Calc\_CGRP\_IAPP; 1.  
 DR PRINTS: PR00817; CALCITONINB.  
 DR PRINTS: PR00818; ISLETAMYLID.  
 DR SMART: SM00113; CALCITONIN; 1.  
 DR PROSITE: PS00258; CALCITONIN; 1.  
 KW AmiAdacton; Amyloid; Hormone.  
 FT MOD\_RES 37 37  
 FT DISULFID 2 7  
 FT SEQUENCE 37 AA; 3921 MW; FE433D9905EBF82E CRC64;  
 Query Match 91.1%; Score 184; DB 1; Length 37;  
 Best Local Similarity 89.2%; Pred. No. 2e-18;  
 Matches 33; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVRSNNLGPVLPPTVNGSNTY 37  
 Db 1 KCNTATCATQRLANFLVRSNNLGPVLPPTVNGSNTY 37  
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 IAPP MESAU STANDARD; PRT; 92 AA.  
 AC P23442;  
 DT 01-NOV-1991 (Rel. 20, Created)  
 DT 01-NOV-1991 (Rel. 20, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Islet amyloid polypeptide precursor (Amylin).  
 GN Name=IAPP;  
 OS Mesocricetus auratus (Golden hamster).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

OC Muroidea; Cricetidae; Cricetinae; Mesocricetus.  
 OX NCBI\_TaxId=10036;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=91067499; PubMed=2251153;  
 RA Nishi M., Bell G.I., Steiner D.F.;  
 RT "Sequence of a cDNA encoding Syrian hamster islet amyloid polypeptide precursor."  
 RL Nucleic Acids Res. 18:6726-6726(1990).  
 CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose utilization and glycogen deposition in muscle, while not affecting adipocyte glucose metabolism.  
 CC -!- SUBCELLULAR LOCATION: Secreted.  
 CC -!- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
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 CC -----  
 CC EMBL: X56067; CAA39545.1; -; mRNA.  
 CC PIR: S13116; S13116.  
 DR InterPro: IPR000443; Amylin.  
 DR InterPro: IPR001693; Calcitonin-like.  
 DR InterPro: IPR002163; Calcitonin B.  
 DR Pfam: PF00214; Calc CGRP IAPP; 1.  
 DR PRINTS: PR00817; CALCITONINB.  
 DR PRINTS: PR00818; ISLETAMYLID.  
 DR SMART: SM00113; CALCITONIN; 1.  
 DR PROSITE: PS00258; CALCITONIN; 1.  
 KW Amidation; Amyloid; Cleavage on pair of basic residues; Hormone; Signal.  
 FT SIGNAL 1 22 Potential.  
 FT PROPEP 23 34 Islet amyloid polypeptide.  
 FT PROPEP 77 92 Tyrosine amide (G-74 provides amide group).  
 FT MOD\_RES 73 73 By similarity.  
 FT DISULFID 38 43  
 FT SEQUENCE 92 AA; 9899 MW; 6D2F7359C4A1D029 CRC64;  
 SQ  
 Query Match 91.1%; Score 184; DB 1; Length 92;  
 Best Local Similarity 89.2%; Pred. 5.7e-18;  
 Matches 33; Conservative 2; Mismatches 2; Indels 0; Gaps 0;  
 QY 1 KONTATCATQRLANFLVHSSNFGPILPTNGSNTY 37  
 DB 37 KONTATCATQRLANFLVHSSNFGPILPTNGSNTY 73  
 RESULT 5  
 IAPP HUMAN  
 ID IAPP\_HUMAN STANDARD; PRT; 89 AA.  
 AC P10997; Q14598;  
 DT 01-JUL-1989 (Rel. 11, Created)  
 DT 01-JUL-1989 (Rel. 11, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Islet amyloid polypeptide precursor (Diabetes-associated peptide) (DAP) (Amylin) (Insulinoma amyloid peptide).  
 GN Name=IAPP;  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homnidae; Homo.  
 OC NCBI\_TaxId=9606;  
 OX  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=8921134; PubMed=2651160; DOI=10.1016/0014-5793(89)81260-8;  
 RA Mosselman S., Hoepfner J.W.M., Lips C.J.M., Janz H.S.;  
 RT "The complete islet amyloid polypeptide precursor is encoded by two exons."  
 RL Proc. Natl. Acad. Sci. U.S.A. 86:154-158(1989).  
 RL

RN [2]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=90114181; PubMed=2608057;  
 RA Nishi M., Sanke T., Seino S., Eddy R.L., Fan Y.-S., Byers M.G., Shows T.B., Bell G.I., Steiner D.F.;  
 RT "Human islet amyloid polypeptide gene: complete nucleotide sequence, chromosomal localization, and evolutionary history."  
 RL Mol. Endocrinol. 3:1775-1781(1989).  
 RN [3]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=89034238; PubMed=3053705;  
 RA Sanke T., Bell G.I., Sample C., Rubenstein A.H., Steiner D.F.;  
 RT "An islet amyloid peptide is derived from an 89-amino acid precursor by proteolytic processing."  
 RL J. Biol. Chem. 263:17243-17246(1988).  
 RN [4]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=90306394; PubMed=2365085; DOI=10.1016/0014-5793(90)80314-9;  
 RA Christman L., Rorsman F., Stenman G., Westermark P., Betsholtz C.;  
 RT "The human islet amyloid polypeptide (IAPP) gene. Organization, chromosomal localization and functional identification of a promoter region."  
 RL PNAS Lett. 267:160-166(1990).  
 RN [5]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=91027936; PubMed=2223985; DOI=10.1016/0167-4781(90)90210-S;  
 RA van Mansfeld A.D.M., Mosselman S., Hoepfner J.W.M., Zandberg J., van Teeffelen H.A.A.M., Baas P.D., Lips C.J.M., Janz H.S.;  
 RT "Islet amyloid polypeptide: structure and upstream sequences of the IAPP gene in rat and man."  
 RL Biochim. Biophys. Acta 1087:235-240(1990).  
 RN [6]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=93129228; PubMed=1282806;  
 RA Hoepfner J.W.M., Oosterwijk C., Visser-Vernooij H.J., Lips C.J.M., Janz H.S.;  
 RT "Characterization of the human islet amyloid polypeptide/amylin gene transcripts: identification of a new polyadenylation site."  
 RL Biochem. Biophys. Res. Commun. 189:1569-1577(1992).  
 RN [7]  
 RP NUCLEOTIDE SEQUENCE OF 28-89.  
 RX MEDLINE=89031237; PubMed=3181427; DOI=10.1016/0014-5793(88)80922-0;  
 RA Mosselman S., Hoepfner J.W.M., Zandberg J., van Mansfeld A.D.M., Geurts van Kessel A.H.M., Lips C.J.M., Janz H.S.;  
 RT "Islet amyloid polypeptide: identification and chromosomal localization of the human gene."  
 RL PNAS Lett. 239:227-232(1988).  
 RN [8]  
 RP PROTEIN SEQUENCE OF 34-52.  
 RX MEDLINE=87048863; PubMed=3535798;  
 RA Westermark P., Wernstedt C., Wilander E., Sletten K.;  
 RT "A novel peptide in the calcitonin gene related peptide family as an amyloid fibril protein in the endocrine pancreas."  
 RL Biochem. Biophys. Res. Commun. 140:827-831(1986).  
 RN [9]  
 RP PROTEIN SEQUENCE OF 34-70.  
 RX MEDLINE=87231921; PubMed=3035556;  
 RA Westermark P., Wernstedt C., Wilander E., Hayden D.W., O'Brien T.D., Johnson K.H.;  
 RT "Amyloid fibrils in human insulinoma and islets of Langerhans of the diabetic cat are derived from a neuropeptide-like protein also present in normal islet cells."  
 RL Proc. Natl. Acad. Sci. U.S.A. 84:3881-3885(1987).  
 RN [10]  
 RP PROTEIN SEQUENCE OF 30-89.  
 RX MEDLINE=9009324; PubMed=2690069;  
 RA Roberts A.N., Leighton B., Todd J.A., Cockburn D., Schofield P.N., Sutton R., Holt S., Boyd Y., Day A.J., Foot E.A., Willis A.C., Reid K.B.M., Cooper G.J.S.;  
 RT "Molecular and functional characterization of amylin, a peptide associated with type 2 diabetes mellitus."  
 RL Proc. Natl. Acad. Sci. U.S.A. 86:9662-9666(1989).  
 RN [11]





SO SEQUENCE 60 AA; 6485 MW; 5DBA512AE8994AE CRC64;  
Query Match 39.1%; Score 79; DB 2; Length 60;  
Best Local Similarity 53.1%; Pred. No. 0.0024;  
Matches 17; Conservative 1; Mismatches 14; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVG 33  
DB 28 CNTATCVTHRLADLRSRGVKSXNFVPTNVG 59  
RESULT 46  
O9GLK2\_RABIT  
ID O9GLK2\_RABIT PRELIMINARY; PRT; 67 AA.  
AC O9GLK2;  
DT 01-MAR-2001 (TrEMBLrel. 16, Created)  
DT 01-MAR-2001 (TrEMBLrel. 16, Last sequence update)  
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)  
DE Calcitonin gene-related peptide variant 1 (Fragment).  
OS Oryctolagus cuniculus (Rabbit).  
OC Eukaryota; Metazoa; Chordata; Craniala; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporidae;  
OC Oryctolagus.  
OC NCBI\_TaxID=9986;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RA Davis M.C., Gierach W.W., Russo A.F.;  
RL Submitted (APR-2000) to the EMBL/GenBank/DBJ databases.  
EMBL; AF260272; AGL1553.1; -; mRNA.  
DR GO; GO:0005576; Cerebellar region; IEA.  
DR GO; GO:0005179; F-hormone activity; IEA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR InterPro; IPR002163; Calcitonin-B.  
DR Pfam; PF00214; Calc CGRP IAPP; I.  
DR PRINTS; PR00817; CALCITONINB.  
DR PRINTS; PR00818; ISLETAMYLID.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; 1.  
FT NON TER 1 1  
FT NON TER 67 67  
SQ SEQUENCE 67 AA; 7286 MW; 99CC326C159BC501 CRC64;  
Query Match 39.1%; Score 79; DB 2; Length 67;  
Best Local Similarity 53.1%; Pred. No. 0.0027;  
Matches 17; Conservative 0; Mismatches 15; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVG 33  
DB 35 CNTATCVTHRLADLRSRGVKSXNFVPTNVG 66  
RESULT 47  
O75V95\_BOVIN  
ID O75V95\_BOVIN PRELIMINARY; PRT; 125 AA.  
AC O75V95;  
DT 05-JUL-2004 (TrEMBLrel. 27, Created)  
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)  
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)  
DE Calcitonin receptor-stimulating peptide-1.  
OS Bos taurus (Bovine).  
OC Eukaryota; Metazoa; Chordata; Craniala; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;  
OC Pecora; Bovidae; Bovinae; Bos.  
OC NCBI\_TaxID=9913;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RA PubMed=14672700; DOI=10.1016/j.birc.2003.11.114;  
RA Katafuchi T., Hamano K., Minamino N.;  
RT Identification, structural determination, and biological activity of  
RT bovine and canine calcitonin receptor-stimulating peptides.;  
RL Biochem. Biophys. Res. Commun. 313:74-79(2004).  
DR EMBL; AB125101; BAD05115.1; -; mRNA.

DR GO; GO:0005576; Cerebellar region; IEA.  
DR GO; GO:0005179; F-hormone activity; IEA.  
DR GO; GO:0004872; Pituitary activity; IEA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR InterPro; IPR002163; Calcitonin-B.  
DR Pfam; PF00214; Calc CGRP IAPP; I.  
DR PRINTS; PR00817; CALCITONINB.  
DR PRINTS; PR00818; ISLETAMYLID.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; UNKNOWN\_1.  
KW RECEPTOR.  
SQ SEQUENCE 125 AA; 14356 MW; 39929D89FB61CFD7 CRC64;  
Query Match 36.6%; Score 74; DB 2; Length 125;  
Best Local Similarity 50.0%; Pred. No. 0.028;  
Matches 16; Conservative 3; Mismatches 13; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVG 33  
DB 81 CNTATCVTHRLADLRSRGVKSXNFVPTNVG 112  
RESULT 48  
Q766Y6\_PIG  
ID Q766Y6\_PIG PRELIMINARY; PRT; 125 AA.  
AC Q766Y6;  
DT 05-JUL-2004 (TrEMBLrel. 27, Created)  
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)  
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)  
DE Calcitonin receptor-stimulating peptide-3.  
OS Sus scrofa (Pig).  
OC Eukaryota; Metazoa; Chordata; Craniala; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suidae;  
OC Sus.  
OC NCBI\_TaxID=9823;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RA MEDLINE=22796190; PubMed=12914769; DOI=10.1016/S0006-291X(03)01413-X;  
RA Katafuchi T., Hamano K., Kikumoto K., Minamino N.;  
RT Identification of second and third calcitonin receptor-stimulating  
RT peptides in porcine brain.;  
RL Biochem. Biophys. Res. Commun. 308:445-451(2003).  
EMBL; AB114134; BAC81766.1; -; mRNA.  
DR GO; GO:0005576; Cerebellar region; IEA.  
DR GO; GO:0005179; F-hormone activity; IEA.  
DR GO; GO:0004872; Pituitary activity; IEA.  
DR InterPro; IPR001693; Calcitonin-like.  
DR InterPro; IPR002163; Calcitonin-B.  
DR Pfam; PF00214; Calc CGRP IAPP; I.  
DR PRINTS; PR00817; CALCITONINB.  
DR SMART; SM00113; CALCITONIN; 1.  
KW RECEPTOR.  
SQ SEQUENCE 125 AA; 14036 MW; 6FE166F9D8907F8E CRC64;  
Query Match 35.1%; Score 71; DB 2; Length 125;  
Best Local Similarity 45.9%; Pred. No. 0.073;  
Matches 17; Conservative 4; Mismatches 8; Indels 8; Gaps 2;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVG 34  
DB 81 CNTATCVTHRLADLRSRGVKSXNFVPTNVG 113  
RESULT 49  
Q862B1\_PIG  
ID Q862B1\_PIG PRELIMINARY; PRT; 126 AA.  
AC Q862B1;  
DT 01-JUN-2003 (TrEMBLrel. 24, Created)  
DT 01-JUN-2003 (TrEMBLrel. 24, Last sequence update)  
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)  
DE Calcitonin receptor-stimulating peptide.  
GN Name=CRSP;

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FT MOD_RES 37 37 Phenylalanine amide.
RL DISTLFRD 2 7 By similarity.
SQ SEQUENCE 37 AA; 3814 MW; 04CDB8BD48B64BA CRC64;

Query Match
Best Local Similarity 40.1%; Score 81; DB 1; Length 37;
Matches 17; Conservative 2; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANPLVHSSNNGPILPPPTNGSNTY 37
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 2 CNTATCVTHRLAGLLSRSGGVKSNFVPTDVGSEAF 37

RESULT 43
Q8WXX2_CALJA PRELIMINARY; PRT; 53 AA.
AC Q8WXX2;
DT 01-MAR-2002 (TREMBLrel. 20, Created)
DT 01-MAR-2002 (TREMBLrel. 20, Last sequence update)
DT 01-JUN-2003 (TREMBLrel. 24, Last annotation update)
DE Beta-calliconin-related protein (Fragment).
OS Name=Beta-CGRP;
GN Callithrix jacchus (Common marmoset).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Primates; Platyrrhini;
OC Callitrichidae; Callitrich.
NCBI_Taxid=9483;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Schindler M., Fischer E.;
RL Submitted (OCT-2001) to the EMBL/Genbank/DBJ databases.
DR EMBL; AF442154; AAL35593.1; -; Genomic DNA.
DR GO; GO:0005576; C:extracellular region; IEA.
DR GO; GO:0005179; F:hormone activity; IEA.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin B.
DR Pfam; PF00214; Calc CGRP IAPP; I.
DR PRINTS; PR00817; CALCITONINB.
DR PRINTS; PR00818; ISLETAMYLOID.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR NON_TER 1
SQ SEQUENCE 53 AA; 5710 MW; A76A34E3D5E3E99C CRC64;

Query Match
Best Local Similarity 40.1%; Score 81; DB 2; Length 53;
Matches 17; Conservative 2; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANPLVHSSNNGPILPPPTNGSNTY 37
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 9 CNTATCVTHRLAGLLSRSGGVKSNFVPTDVGSKAF 44

RESULT 44
CALCB_HORSE STANDARD; PRT; 129 AA.
ID CALCB_HORSE
AC Q9N0T3;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Calcitonin gene-related peptide II precursor (GGRP-II) (beta-type
DE CGRP)
OS Name=CALCB;
OS Equus caballus (Horse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Perissodactyla; Equidae; Equus.
NCBI_Taxid=9796;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA MEDLINE=22470155; PubMed=12581884; DOI=10.1016/S0303-7207(02)00289-7;
RA Toribio R.E., Kohn C.W., Leone G.W., Capen C.C., Rosol T.J.;
RA "Molecular cloning and expression of equine calcitonin, calcitonin
```

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RL gene-related peptide-I, and calcitonin gene-related peptide-II.";
RL Mol. Cell. Endocrinol. 199;119-128(2003).
CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of
CC vessels including the coronary, cerebral and systemic vasculature.
CC Its abundance in the CNS also points toward a neurotransmitter or
CC neuromodulator role (By similarity).
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- SIMILARITY: Belongs to the calcitonin family.
CC
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CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC the European Bioinformatics Institute. There are no restrictions on its
CC use as long as its content is in no way modified and this statement is not
CC removed.
CC -----
CC EMBL; AF257470; AAF70199.1; -; mRNA.
CC InterPro; IPR000443; Amylin.
CC InterPro; IPR001693; Calcitonin-like.
CC InterPro; IPR002163; Calcitonin B.
CC Pfam; PF00214; Calc CGRP IAPP; I.
CC PRINTS; PR00817; CALCITONINB.
CC PRINTS; PR00818; ISLETAMYLOID.
CC SMART; SM00113; CALCITONIN; 1.
CC PROSITE; PS00258; CALCITONIN; 1.
KW Amidation; Cleavage on pair of basic residues; Hormone; Signal.
FT SIGNAL 1 25 Potential.
FT PROPEP 26 81 By similarity.
FT PEPTIDE 84 120 Calcitonin gene-related peptide II.
FT PROPEP 126 129 By similarity.
FT MOD_RES 120 129 Phenylalanine amide (G-121 provides amide
FT group) (By similarity).
FT DISTLFRD 85 90 By similarity.
SQ SEQUENCE 129 AA; 13925 MW; 97C2C7AC713ABD9B CRC64;

Query Match
Best Local Similarity 39.6%; Score 80; DB 1; Length 129;
Matches 17; Conservative 47.2%; Pred. No. 0.0041; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANPLVHSSNNGPILPPPTNGSNTY 37
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DB 85 CNTATCVTHRLAGLLSRSGGVKSNFVPTDVGSEAF 120

RESULT 45
Q9GLK1_RABIT PRELIMINARY; PRT; 60 AA.
ID Q9GLK1;
DT 01-MAR-2001 (TREMBLrel. 16, Created)
DT 01-MAR-2001 (TREMBLrel. 16, Last sequence update)
DT 01-JUN-2003 (TREMBLrel. 24, Last annotation update)
DE Calcitonin gene-related peptide (Fragment).
OS Oryctolagus cuniculus (Rabbit).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporidae;
OC Oryctolagus.
NCBI_Taxid=9986;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Davis M.C., Gierasch W.W., Russo A.F.;
RL Submitted (APR-2000) to the EMBL/Genbank/DBJ databases.
DR EMBL; AF260273; AAG15536.1; -; mRNA.
DR GO; GO:0005576; C:extracellular region; IEA.
DR GO; GO:0005179; F:hormone activity; IEA.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin B.
DR Pfam; PF00214; Calc CGRP IAPP; I.
DR PRINTS; PR00817; CALCITONINB.
DR PRINTS; PR00818; ISLETAMYLOID.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR NON_TER 1
FT SEQUENCE 60 60
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CC use as long as its content is in no way modified and this statement is not  
CC removed.  
-----  
DR EMBL: M1596; AAA40850.1; -; mRNA.  
DR PIR: A44173; A44173.  
DR Ensembl: ENSRNOC0000011074; Rattus norvegicus.  
DR RGD: 620997; CalcB.  
DR InterPro: IPR000443; Amylin.  
DR InterPro: IPR001693; Calcitonin-like.  
DR InterPro: IPR002163; Calcitonin B.  
DR Pfam: PF00214; Calc CGRP IAPP; I.  
DR PRINTS: PR00817; CALCITONINB.  
DR SMART: SM00113; CALCITONIN.  
DR PROSITE: PS00258; CALCITONIN; 1.  
KM Amidation; Cleavage on pair of basic residues; Hormone; Signal.  
FT SIGNAL 1 26  
FT PROPEP 27 86 By similarity.  
FT PEPTIDE 89 125 Calcitonin gene-related peptide II.  
FT PROPEP 131 134 By similarity.  
FT MOD\_RES 125 125 Phenylalanine amide (G-126 provides amide  
FT DISULFID 90 95 group) (By similarity).  
SQ SEQUENCE 134 AA; 14965 MW; BF6CAF8B74489B38 CRC64;  
  
Query Match 41.6%; Score 84; DB 1; Length 134;  
Best Local Similarity 50.0%; Pred. No. 0.0012;  
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;  
  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVSNTY 37  
Db 90 CNTATCVTHRLAGLRRSGGVTKDNFVPTNVSKEAF 125  
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RESULT 41  
CALCB\_MOUSE STANDARD; PRT; 130 AA.  
ID CALCB\_MOUSE  
AC 099MP3;  
DT 28-FEB-2003 (Rel. 41, Created)  
DT 28-FEB-2003 (Rel. 41, Last sequence update)  
DT 10-MAY-2005 (Rel. 47, Last annotation update)  
DE Calcitonin gene-related peptide II precursor (CGRP-II) (Beta-type  
DE CGRP).  
GN Name=CalcB;  
OS Mus musculus (Mouse).  
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;  
OC Muridae; Muridae; Murinae; Mus.  
OX NCBI\_TaxID=10090;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RC STRAIN=129/Sv.  
RX MEDLINE=21604266; PubMed=11761712;  
RA Thomas P.M., Nasonkin I., Zhang H., Gage R.F., Cole G.J.;  
RT "Structure of the mouse calcitonin/calcitonin gene-related peptide  
RT alpha and beta genes.";  
RL DNA Seq. 12:131-135(2001).  
CC -!- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
CC vessels including the coronary, cerebral and systemic vasculature.  
CC Its abundance in the CNS also points toward a neurotransmitter or  
CC neuromodulator role (By similarity).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- SIMILARITY: Belongs to the calcitonin family.  
-----  
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CC use as long as its content is in no way modified and this statement is not  
CC removed.  
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DR EMBL: AF325526; AAK16431.1; -; Genomic DNA.  
DR EMBL: AF325524; AAK16431.1; JOINED; Genomic DNA.  
DR Ensembl: ENSMUSG0000030666; Mus musculus.  
DR MGI: MGI:2151254; CalcB.  
DR GO: GO:0005615; Extracellular space; TAS.  
DR InterPro: IPR000443; Amylin.  
DR InterPro: IPR001693; Calcitonin-like.  
DR InterPro: IPR002163; Calcitonin B.  
DR Pfam: PF00214; Calc CGRP IAPP; I.  
DR PRINTS: PR00817; CALCITONINB.  
DR SMART: SM00113; CALCITONIN.  
DR PROSITE: PS00258; CALCITONIN; 1.  
KM Amidation; Cleavage on pair of basic residues; Hormone; Signal.  
FT SIGNAL 1 26  
FT PROPEP 27 82 By similarity.  
FT PEPTIDE 84 120 Calcitonin gene-related peptide II.  
FT PROPEP 127 130 By similarity.  
FT MOD\_RES 120 120 Phenylalanine amide (G-121 provides amide  
FT DISULFID 85 90 group) (By similarity).  
SQ SEQUENCE 130 AA; 14623 MW; 97299244B8F6C36 CRC64;  
  
Query Match 40.6%; Score 82; DB 1; Length 130;  
Best Local Similarity 47.2%; Pred. No. 0.0021;  
Matches 17; Conservative 3; Mismatches 16; Indels 0; Gaps 0;  
  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVSNTY 37  
Db 85 CNTATCVTHRLADLSRSGGVTKDNFVPTNVSKEAF 120  
-----  
RESULT 42  
CALCA\_PIG STANDARD; PRT; 37 AA.  
ID CALCA\_PIG  
AC P30860;  
DT 01-JUL-1993 (Rel. 26, Created)  
DT 01-JUL-1993 (Rel. 26, Last sequence update)  
DT 10-MAY-2005 (Rel. 47, Last annotation update)  
DE Calcitonin gene-related peptide (CGRP).  
GN Name=CALCA; Synonyms=CALC;  
OS Sus scrofa (Pig).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suidae;  
OC Sus.  
OX NCBI\_TaxID=9823;  
RN [1]  
RP PROTEIN SEQUENCE.  
RX MEDLINE=87173481; PubMed=3494209; DOI=10.1016/0143-4179(87)90034-5;  
RA Kimura S., Sugita Y., Kanazawa I., Saito A., Goto K.;  
RT "Isolation and amino acid sequence of calcitonin gene related peptide  
RT from porcine spinal cord.";  
RL Neuropeptides 9:75-82(1987).  
CC -!- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
CC vessels including the coronary, cerebral and systemic vasculature.  
CC Its abundance in the CNS also points toward a neurotransmitter or  
CC neuromodulator role.  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- SIMILARITY: Belongs to the calcitonin family.  
-----  
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CC between the Swiss Institute of Bioinformatics and the EMBL Outstation -  
CC the European Bioinformatics Institute. There are no restrictions on its  
CC use as long as its content is in no way modified and this statement is not  
CC removed.  
-----  
DR InterPro: IPR001693; Calcitonin-like.  
DR InterPro: IPR002163; Calcitonin B.  
DR Pfam: PF00214; Calc CGRP IAPP; I.  
DR PRINTS: PR00817; CALCITONINB.  
DR SMART: SM00113; CALCITONIN.  
DR PROSITE: PS00258; CALCITONIN; 1.  
KM Amidation; Direct protein sequencing; Hormone.

RX MEDLINE=87713163; PubMed=3034287;  
 RA Craig R.K., Riley J.H., Edbrooke M.R., Broad P.M., Foord S.M.,  
 RA Al-Kazwini S.J., Holman J.J., Marshall I.,  
 RT "Expression and function of the human calcitonin/alpha-CGRP gene in  
 RT health and disease.";  
 RL Biochem. Soc. Symp. 52:91-105(1986).  
 RN [7]  
 RP PROTEIN SEQUENCE OF 83-119.  
 RX MEDLINE=84191466; PubMed=6609312;  
 RA Morris H.R., Panico M., Etienne T., Tipping J., Girgis S.I.,  
 RA McIntyre I.,  
 RT "Isolation and characterization of human calcitonin gene-related  
 RT peptide.";  
 RL Nature 308:746-748(1984).  
 RN [8]  
 RP PARTIAL PROTEIN SEQUENCE OF 83-117.  
 RX MEDLINE=87109142; PubMed=3492492;  
 RA Petermann J.B., Born W., Chang J.Y., Fischer J.A.,  
 RT "Identification in the human central nervous system, pituitary, and  
 RT thyroid of a novel calcitonin gene-related peptide, and partial amino  
 RT acid sequence in the spinal cord.";  
 RL J. Biol. Chem. 262:542-545(1987).  
 RN [9]  
 RP PROTEIN SEQUENCE OF 83-108, AND FUNCTION.  
 RC TISSUE=phoechromocytoma;  
 RX MEDLINE=92287083; PubMed=1318039;  
 RA Kitamura K., Kangawa K., Kawamoto M., Ichiki Y., Matsuo H., Eto T.,  
 RT "Isolation and characterization of peptides which act on rat  
 RT platelets, from a phoechromocytoma.";  
 RL Biochem. Biophys. Res. Commun. 185:134-141(1992).  
 RN [10]  
 RP STRUCTURE BY NMR OF CGRP.  
 RX MEDLINE=91105142; PubMed=1988044;  
 RA Breese A.L., Harvey T.S., Bazzo R., Campbell I.D.,  
 RT "Solution structure of human calcitonin gene-related peptide by 1H NMR  
 RT and distance geometry with restrained molecular dynamics.";  
 RL Biochemistry 30:575-582(1991).  
 RN [11]  
 RP STRUCTURE BY NMR OF CGRP.  
 RX MEDLINE=91248117; PubMed=2039456;  
 RA Hubbard J.A.M., Martin S.R., Chaplin L.C., Bose C., Kelly S.M.,  
 RA Price N.C.,  
 RT "Solution structures of calcitonin-gene-related-peptide analogues of  
 RT calcitonin-gene-related peptide and amylin.";  
 RL Biochem. J. 275:785-788(1991).  
 CC -I- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 CC vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 CC neuromodulator role. It also elevates platelet CAMP.  
 CC -I- ALTERNATIVE PRODUCTS:  
 CC Event=Alternative splicing; Named isoforms=3;  
 CC Name=3;  
 CC IsoId=P06881-1; Sequence=Displayed;  
 CC Name=1;  
 CC IsoId=P01258-1; Sequence=External;  
 CC Name=2;  
 CC IsoId=P01258-2; Sequence=External;  
 CC -I- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
 CC This Swiss-Prot entry is copyright. It is produced through a collaboration  
 CC between the Swiss Institute of Bioinformatics and the EMBL outstation -  
 CC the European Bioinformatics Institute. There are no restrictions on its  
 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 CC EMBL; M12667; AAA51914.1; -; Genomic DNA.  
 CC EMBL; M12664; AAA51914.1; JOINED; Genomic DNA.  
 CC EMBL; M12665; AAA51914.1; JOINED; Genomic DNA.  
 CC EMBL; X15943; CAA34070.1; -; Genomic DNA.  
 CC EMBL; K03512; AAA52011.1; -; mRNA.  
 CC EMBL; X02350; CAA26190.1; -; mRNA.  
 CC EMBL; M28637; AAA52012.1; -; Genomic DNA.  
 CC EMBL; M26094; AAA51912.1; -; Genomic DNA.

DR PIR: S07644; TCHUR.  
 DR PDB: 1LS7; Model: A=1-125.  
 DR Ensembl: ENSG00000110680; Homo sapiens.  
 DR HGNC: HGNC:1437; CALCA.  
 DR MIM: 114130; -;  
 DR GO: GO:0005783; C: endoplasmic reticulum; TAS.  
 DR GO: GO:0005615; C: extracellular space; TAS.  
 DR GO: GO:0005625; C: soluble fraction; TAS.  
 DR GO: GO:0005102; F: receptor binding; TAS.  
 DR GO: GO:0007190; P: adenylate cyclase activation; TAS.  
 DR GO: GO:0007267; P: cell-cell signaling; TAS.  
 DR GO: GO:0007188; P: G-protein signaling, coupled to CAMP nucleo. . .; TAS.  
 DR GO: GO:0007202; P: phospholipase C activation; TAS.  
 DR GO: GO:0007204; P: positive regulation of cytosolic calcium io. . .; TAS.  
 DR GO: GO:0008217; P: regulation of blood pressure; TAS.  
 DR GO: GO:0001501; P: skeletal development; TAS.  
 DR InterPro: IPR001693; Calcitonin-like.  
 DR InterPro: IPR002163; Calcitonin B.  
 DR Pfam: PF00214; Calc CGRP APP. I.  
 DR PRINTS: PR00817; CALCITONINB.  
 DR SMART: SM00113; CALCITONIN; 1.  
 DR PROSITE: PS00258; CALCITONIN; 1.  
 KW 3D-structure; Alternative splicing; Amidation;  
 KW Cleavage on pair of basic residues; Direct protein sequencing;  
 KW Hormone; Signal.  
 FT SIGNAL 1 25 Potential.  
 FT PROPEP 26 80  
 FT PEPPTIDR 83 119 Calcitonin gene-related peptide I.  
 FT PROPEP 125 128  
 FT MOD\_RES 119 119 Phenylalanine amide (G-120 provides amide  
 FT group).  
 FT DISULFID 84 89  
 SQ SEQUENCE 128 AA; 1389 MW; A003A1069260D988 CRC64;  
 Query Match 41.6%; Score 84; DB 1; Length 128;  
 Best Local Similarity 50.0%; Pred. No. 0.0011;  
 Matches 20; Conservative 2; Mismatches 10; Indels 8; Gaps 2;  
 OY 2 CNTATCATQRIANFLVHS---SNNFGRLPPVNGSNTY 37  
 Db 84 CDTATCVTHRLAGLLSRSGGVKNF---VPTNVGSKAF 119  
 RESULT 40  
 CALCB\_RAT STANDARD; PRT; 134 AA.  
 ID CALCB\_RAT  
 AC P10093;  
 DT 01-MAR-1989 (Rel. 10, Created)  
 DT 01-MAR-1989 (Rel. 10, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DB Calcitonin gene-related peptide II precursor (CGRP-II) (Beta-type  
 DB CGRP).  
 GN Name=Calcb;  
 OS Rattus norvegicus (Rat).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;  
 OC Muridae; Muridae; Murinae; Rattus.  
 OX NCBI\_TaxID=10116;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=85300490; PubMed=2994212;  
 RA Amara S.G., Arizze J.L., Leff S.E., Swanson L.W., Evans R.M.,  
 RA Rosenfeld M.G.,  
 RT "Expression in brain of a messenger RNA encoding a novel neuropeptide  
 RT homologous to calcitonin gene-related peptide.";  
 RL Science 229:1094-1097(1985).  
 CC -I- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 CC vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 CC neuromodulator role.  
 CC -I- SUBCELLULAR LOCATION: Secreted.  
 CC -I- SIMILARITY: Belongs to the calcitonin family.

DB 84 CNTATCVTHRLAGLSRSGGVKDNFVPTNVGSEAF 119

RESULT 37  
O8WNX3 CALJA  
ID O8WNX3 CALJA PRELIMINARY; PRT; 52 AA.

DT 01-MAR-2002 (TrEMBLrel. 20, Created)  
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)  
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)  
DE Alpha-calcitonin-related protein (Fragment).  
GN Name=alpha-CGRP;  
OS Callitrix jacchus (Common marmoset).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Euarchontoglires; Primates; Platyrrhini;  
OC Callitrichidae; Callitrichi.  
OX NCBI\_TaxId=9483;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RA Schindler M., Fischer E.;  
RL Submitted (OCT-2001) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AF442153; AAL35592.1; -; Genomic DNA.  
DR GO; GO:0005576; C:extracellular region; IEA.  
DR GO; GO:0005179; F:hormone activity; IEA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR pfam; PF00214; Calc CGRP IAPP; 1.  
DR PRINTS; PR00817; CALCITONIN.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; 1.  
SQ  
NON\_TER 1  
SEQUENCE 52 AA; 5643 MW; 7C0A3ACE63CC4DB1 CRC64;

Query Match 41.6%; Score 84; DB 2; Length 52;  
Best Local Similarity 50.0%; Pred. No. 0.00039;  
Matches 20; Conservative 2; Mismatches 10; Indels 8; Gaps 2;

OY 2 CNTATCATQRIANFLVHS-----SNNFGPILPPTNVGSNTY 37  
DB 8 CDTATCVTHRLAGLSRSGGVKDNF-----VPTNVGSEAF 43

RESULT 38  
O90YC3 PAROL  
ID O90YC3 PAROL PRELIMINARY; PRT; 53 AA.

DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)  
DE Calcitonin gene-related peptide (Fragment).  
GN Name=CT/CGRP;  
OS Paralicthys olivaceus (Japanese flounder).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;  
OC Pleuronectidae; Paralicthyidae; Paralicthys.  
OX NCBI\_TaxId=8255;  
RN [1]  
RP NUCLEOTIDE SEQUENCE.  
RA MEDLINE=21406117; PubMed=11514025; DOI=10.1016/S0196-9781(01)00484-3;  
RT Suzuki N., Suzuki T., Kurokawa T.;  
RT "Cloning of a calcitonin gene-related peptide from genomic DNA and its  
RT mRNA expression in flounder, Paralicthys olivaceus";  
RL Peptide 22:1435-1438(2001).  
DR EMBL; AB052782; BAB6411.1; -; Genomic DNA.  
DR GO; GO:0005576; C:extracellular region; IEA.  
DR GO; GO:0005179; F:hormone activity; IEA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR pfam; PF00214; Calc CGRP IAPP; 1.

DR PRINTS; PR00817; CALCITONIN.  
DR PRINTS; PR00818; ISLETAMYLID.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; 1.  
FT CHAIN 8 44 calcitonin gene-related peptide.  
FT NON\_TER 1 1  
SQ SEQUENCE 53 AA; 5684 MW; 14C0191868A64CF4 CRC64;

Query Match 41.6%; Score 84; DB 2; Length 53;  
Best Local Similarity 47.5%; Pred. No. 0.0004;  
Matches 19; Conservative 4; Mismatches 9; Indels 8; Gaps 2;

OY 2 CNTATCATQRIANFLVHS-----SNNFGPILPPTNVGSNTY 37  
DB 9 CNTSTCVTHRLADLSRSGGLGYNNF-----VPTNVGAQAF 44

RESULT 39  
ID CALCA HUMAN STANDARD; PRT; 128 AA.

AC P06881; O93048; O9UCP0;  
DT 01-JAN-1988 (Rel. 06, Created)  
DT 01-MAR-1989 (Rel. 10, Last sequence update)  
DT 10-MAY-2005 (Rel. 47, Last annotation update)  
DE Calcitonin gene-related peptide I precursor (CGRP-I) (Alpha-type  
DE CGRP).  
GN Name=CALCA; Synonyms=CALC1;  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homiidae;  
OC Homo.  
OX NCBI\_TaxId=9606;  
RN [1]  
RP NUCLEOTIDE SEQUENCE [GENOMIC DNA].  
RX MEDLINE=85166259; PubMed=3872459;  
RA Jones V., Lin C.R., Kawashima E., Semon D., Swanson L.W.,  
RA Mermod J.-D., Evans R.M., Rosenfeld M.G.;  
RT "Alternative RNA processing events in human calcitonin/calcitonin  
RT gene-related peptide gene expression";  
RL Proc. Natl. Acad. Sci. U.S.A. 82:1994-1998(1985).  
RN [2]  
RP NUCLEOTIDE SEQUENCE [GENOMIC DNA].  
RX MEDLINE=89386053; PubMed=2571128;  
RA Broad P.M., Symes A.J., Thakker R.V., Craig R.K.;  
RL "Structure and methylation of the human calcitonin/alpha-CGRP gene";  
RN Nucleic Acids Res. 17:6999-7011(1989).  
RN [3]  
RP NUCLEOTIDE SEQUENCE [MRNA] OF 48-119.  
RX MEDLINE=85022523; PubMed=6148938;  
RA Nelkin B.D., Rosenfeld K.I., de Bustros A., Leong S.S., Roos B.A.,  
RA Baylin S.B.;  
RT "Structure and expression of a gene encoding human calcitonin and  
RT calcitonin gene related peptide";  
RL Biochem. Biophys. Res. Commun. 123:648-655(1984).  
RN [4]  
RP NUCLEOTIDE SEQUENCE [MRNA] OF 49-128.  
RX MEDLINE=85230541; PubMed=2400883;  
RA Edbrooke M.R., Parker D., Mcvey J.H., Riley J.H., Sorenson G.D.,  
RA Petrengill O.S., Craig R.K.;  
RT "Expression of the human calcitonin/CGRP gene in lung and thyroid  
RT carcinoma";  
RL EMBO J. 4:715-724(1985).  
RN [5]  
RP NUCLEOTIDE SEQUENCE [GENOMIC DNA] OF 77-128.  
RC TISSUE=Thyroid carcinoma;  
RX MEDLINE=84240176; PubMed=6610687;  
RA Stenbergh P.H., Hoppener J.W., Zandberg J., de Ven W.J., Jansz H.S.,  
RA Lips C.J.;  
RT "Calcitonin gene related peptide coding sequence is conserved in the  
RT human genome and is expressed in medullary thyroid carcinoma";  
RL J. Clin. Endocrinol. Metab. 59:358-360(1984).  
RN [6]  
RP NUCLEOTIDE SEQUENCE [GENOMIC DNA] OF 77-128.

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RT "Alternative RNA processing events in human calcitonin/calcitonin
RT gene-related peptide gene expression." ;
RL Proc. Natl. Acad. Sci. U.S.A. 82:1994-1998(1985).
RN [2]
RN NUCLEOTIDE SEQUENCE.
RP MEDLINE=86220111; PubMed=6283379;
RA Amara S.G., Jones V., Rosenfeld M.G., Ong E.S., Evans R.M.;
RT "Alternative RNA processing in calcitonin gene expression generates
RT mRNAs encoding different polypeptide products." ;
RL Nature 298:240-244(1982).
RN [3]
RN NUCLEOTIDE SEQUENCE.
RP MEDLINE=85300490; PubMed=2994212;
RA Amara S.G., Arrieta J.L., Leff S.E., Swanson L.W., Evans R.M.,
RA Rosenthal M.G.;
RT "Expression in brain of a messenger RNA encoding a novel neuropeptide
RT homologous to calcitonin gene-related peptide." ;
RL Science 229:1094-1097(1985).
CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of
CC vessels including the coronary, cerebral and systemic vasculature.
CC Its abundance in the CNS also points toward a neurotransmitter or
CC neuromodulator role.
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- ALTERNATIVE PRODUCTS:
CC Event=Alternative splicing; Named isoforms=2;
CC Name=Calcitonin-gene related peptide 1;
CC IsoId=P01256-1; Sequence=Displayed;
CC Name=Calcitonin;
CC IsoId=P01257-1; Sequence=External;
CC -1- SIMILARITY: Belongs to the calcitonin family.
-----
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CC removed.
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DR EMBL; L29188; AAB59682.1; -; Genomic DNA.
DR EMBL; L00109; AAB59682.1; JOINED; Genomic DNA.
DR EMBL; L00110; AAB59682.1; JOINED; Genomic DNA.
DR EMBL; V01231; CAA24541.1; -; mRNA.
DR EMBL; M11597; AAA40847.1; -; mRNA.
DR PIR; A01524; TCRR.
DR PIR; B44173; B44173.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin B.
DR Pfam; PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS; PR00817; CALCITONINB.
DR PRINTS; PR00818; ISLETAMYLOID.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
KW Alternative splicing; Amidation; Cleavage on pair of basic residues;
KW Hormone; Signal.
FT SIGNAL 1 25
FT PROPEP 26 80
FT PEPTIDE 83 119 Calcitonin gene-related peptide 1.
FT PROPEP 125 128
FT MOD_RES 119 119 Phenylalanine amide (G-120 provides amide
FT group).
FT DISULFD 84 89 By similarity.
FT CONFLICT 40 40 Missing (in Ref. 2 and 3).
FT CONFLICT 51 51 Missing (in Ref. 2 and 3).
FT CONFLICT 70 70 Q -> E80 (in Ref. 2 and 3).
FT CONFLICT 99 99 S -> R (in Ref. 3).
SQ SEQUENCE 128 AA; 13948 MW; 75D14869C17078D3 CRC64;
Query Match 42.1%; Score 85; DB 1; Length 128;
Best Local Similarity 50.0%; Pred. No. 0.0008;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0.

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OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
 OC Acanthopterygia; Acanthopterygii; Perciformes; Tetraodontiformes;  
 OC Tetraodontidae; Tetraodontidae; Tetraodon.  
 OC NCBI\_TaxId=99883;  
 [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RA Jallion O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,  
 RA Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,  
 RA Nicoud S., Jaife D., Fisher S., Lutfalla G., Dossat C., Segurens B.,  
 RA Desliya C., Sainoubat M., Levy M., Boudet N., Castellano S.,  
 RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,  
 RA Biemont C., Skalli Z., Catolico L., Poulain J., De Bernardis V.,  
 RA Craud C., Duprat S., Broctier P., Couanceau J.P., Gouzy J.,  
 RA Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,  
 RA Kellis M., Volff J.N., Guigo R., Zody M.C., Mesirov J.,  
 RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,  
 RA Laudet V., Schachter V., Querier F., Saurin W., Scarpelli C.,  
 RA Winkler P., Lander B.S., Weissenbach J., Roest Crolius H.,  
 RA "Genome duplication in the teleost fish Tetraodon nigroviridis reveals  
 RT the early vertebrate proto-karyotype."  
 RL Nature 431:946-957(2004).  
 RN [2]  
 RP NUCLEOTIDE SEQUENCE.  
 RG Genoscope, Whitehead Institute Centre for Genome Research;  
 RL Submitted (FEB-2004) to the EMBL/GenBank/DBJ databases.  
 CC -1- CAUTION: The sequence shown here is derived from an  
 CC EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is  
 CC preliminary data.  
 CC EMBL; CAAB01002800; CAP87593.1; -; Genomic\_DNA.  
 FT NON\_TER 1 1  
 FT 44 44  
 SQ SEQUENCE 44 AA; 4556 MW; EE24CED223FAE96 CRC64;  
 Query Match 42.1%; Score 85; DB 2; Length 44;  
 Best Local Similarity 54.1%; Pred. No. 0.00027;  
 Matches 20; Conservative 3; Mismatches 6; Indels 8; Gaps 2;  
 QY 2 CNTATCATGRLANFLVHSS---NNFGPILPPTNVGS 34  
 ID 066VCI RAT PRELIMINARY; PRT; 50 AA.  
 AC 066VCI;  
 DT 25-OCT-2004 (T-EMBLrel. 28, Created)  
 DT 25-OCT-2004 (T-EMBLrel. 28, Last sequence update)  
 DT 25-OCT-2004 (T-EMBLrel. 28, Last annotation update)  
 DE Alpha-calctonin gene-related peptide (Fragment).  
 OS Rattus norvegicus (Rat).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;  
 OC Muridae; Murinae; Rattus.  
 OC NCBI\_TaxId=10116;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RC STRAIN=Mistar;  
 RA Gerritsaotis I., Garcia del Cano G., Canudas J., Sarasa M.,  
 RA Martinez-Millan L.,  
 RA "Evidence for expression of calcitonin gene-related peptide in  
 RT collicular neurons during postnatal development."  
 RL Submitted (JUL-2004) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; A7702025; AA007931.1; -; mRNA.  
 DR GO; GO:0005576; Cerebellar region; IRA.  
 DR GO; GO:0005179; Hormone activity; IRA.  
 DR InterPro; IPR001693; Calcitonin-like.  
 DR Pfam; PF00214; Calc CGRP IAPP; 1.  
 DR PRINTS; PR00817; CALCITONINB.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.

FT NON\_TER 1 1  
 SQ SEQUENCE 50 AA; 5402 MW; 295BEPF036CF7FA CRC64;  
 Query Match 42.1%; Score 85; DB 2; Length 50;  
 Best Local Similarity 50.0%; Pred. No. 0.00027;  
 Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;  
 QY 2 CNTATCATGRLANFLVHSSNNFGPILPPTNVGSNTY 37  
 ID 066VCI RAT PRELIMINARY; PRT; 128 AA.  
 AC 066VCI;  
 DT 28-FEB-2003 (Rel. 41, Created)  
 DT 28-FEB-2003 (Rel. 41, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Calcitonin gene-related peptide I precursor (CGRP-I) (Alpha-type  
 DE CGRP).  
 GN Name=Calc; Synonyms=Calc;  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;  
 OC Muridae; Murinae; Mus.  
 OC NCBI\_TaxId=10090;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RA Sarasa M., Catalan J., Aramayo J., Sorribas V.,  
 RA "Mouse CGRP precursor is highly homologous to that of the rat."  
 RT Submitted (DEC-2000) to the EMBL/GenBank/DBJ databases.  
 RN [2]  
 RP NUCLEOTIDE SEQUENCE.  
 RC STRAIN=129/Sv;  
 RX MEDLINE=21604266; PubMed=11761712;  
 RA Thomas P.M., Nasoskin I., Zhang H., Gagel R.F., Cote G.J.,  
 RT "Structure of the mouse calcitonin/calcitonin gene-related peptide  
 RT alpha and beta genes."  
 RL DNA Seq. 12:131-135(2001).  
 RN [3]  
 RP NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].  
 RC STRAIN=C57BL/6J; TISSUE=Mammary gland;  
 RX MEDLINE=2388257; PubMed=12477932; DOI=10.1073/pnas.242603899;  
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,  
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,  
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,  
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,  
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,  
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
 RA Brownstein M.J., Ushed T.B., Toshiyuki S., Carninci P., Prange C.,  
 RA Raha S.S., Lochellano N.A., Peters G.J., Abramson R.D., Millar S.J.,  
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunnarsson P.H.,  
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hultk S.W.,  
 RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
 RA Fahey J., Heltan E., Kettelman M., Madan A., Rodriguez S., Sanchez A.,  
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,  
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,  
 RA Buterfield Y.S.N., Krzywinski M.I., Skalska U., Smalins D.E.,  
 RA Scherch A., Schein J.E., Jones S.J.M., Marra M.A.,  
 RT "Generation and initial analysis of more than 15,000 full-length human  
 RT and mouse cDNA sequences."  
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).  
 CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 CC vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 CC neuromodulator role. It also elevates platelet cAMP (by  
 CC similarity).  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- ALTERNATIVE PRODUCTS:  
 CC Event=Alternative splicing, Named isoforms=2;  
 CC Name=Calcitonin-gene related peptide I;



FT DISULFID 83 88 G -> S (In Ref. 2).  
 FT CONFLICT 73 73  
 SQ SEQUENCE 127 AA; 13706 MW; B0A71A063CD5ACE7 CRC64;

Query Match 42.6%; Score 86; DB 1; Length 127;  
 Best Local Similarity 50.0%; Pred. No. 0.00057;  
 Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRIANFLVHSSNNFGPILPPTNVGSNTY 37  
 DB 83 CNTATCVTHRLAGLLSRSGGVKSNFVPTNVGSKAF 118

## RESULT 31

056910 HUMAN PRELIMINARY; PRT; 127 AA.  
 AC 056910;  
 DT 10-MAY-2005 (TREMBlrel. 30, Created)  
 DT 10-MAY-2005 (TREMBlrel. 30, Last sequence update)  
 DT 10-MAY-2005 (TREMBlrel. 30, Last annotation update)  
 DE Calcitonin-related polypeptide, beta.  
 GN Name=CALCB;  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Primates; Catherhini; Homnidae;  
 OC Homo  
 NCBI\_TaxID=9606;

## NCBI\_TaxID=9606;

NUCLEOTIDE SEQUENCE.

RC TISSUE=Brain;  
 RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;  
 RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,  
 Altschul S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,  
 Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,  
 Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,  
 Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
 Brown S.S., Loggellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,  
 Bock S.A., McKernan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,  
 Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,  
 Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
 Fahy J., Helton E., Kettelman M., Madan A., Rodighiero S., Sanchez A.,  
 Whiting M., Madan A., Young A.C., Shcherchenko Y., Bouffard G.G.,  
 Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,  
 Rodriguez A.C., Grimwood J., Schmitt J., Myers R.M.,  
 Butcherfield J.S.N., Krzywinski M.I., Skalska U., Smalins D.E.,  
 Scherch A., Schein J.E., Jones S.J.M., Marra M.A.;  
 "Generation and initial analysis of more than 15,000 full-length human  
 RT and mouse cDNA sequences.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).  
 RN [2]  
 RP NUCLEOTIDE SEQUENCE.  
 RC TISSUE=Brain;  
 RG NIH MGC Project;  
 RL Submitted (Apr-2005) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; BC092468; AAH92468.1; -, mRNA.  
 DR Ensemble; ENSG00000175868; Homo sapiens.  
 DR GO; GO:0005576; C:extracellular region; IEA.  
 DR GO; GO:0005179; F:hormone activity; IEA.  
 DR InterPro; IPR000443; Amylin.  
 DR InterPro; IPR001693; Calcitonin-like.  
 DR InterPro; IPR002163; Calcitonin-B.  
 DR Pfam; PF00214; Calc\_CGRP\_IAPP; I.  
 DR PRINTS; PR00817; CALCITONIN.  
 DR PRINTS; PR00818; ISLETAMYLID.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.  
 SQ SEQUENCE 127 AA; 13706 MW; B0A71A063CD5ACE7 CRC64;

Query Match 42.6%; Score 86; DB 2; Length 127;  
 Best Local Similarity 50.0%; Pred. No. 0.00057;  
 Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRIANFLVHSSNNFGPILPPTNVGSNTY 37  
 DB 83 CNTATCVTHRLAGLLSRSGGVKSNFVPTNVGSKAF 118

## RESULT 32

CALCA SHEEP STANDARD; PRT; 37 AA.  
 ID CALCA\_SHEEP  
 AC P30881;  
 DT 01-JUL-1993 (Rel. 26, Created)  
 DT 01-JUL-1993 (Rel. 26, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Calcitonin gene-related peptide (CGRP).  
 GN Name=CALCA; Synonyms=CALC;  
 OS Ovis aries (Sheep).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;  
 OC Pecora; Bovidae; Caprinae; Ovis.  
 NCBI\_TaxID=9940;

## NCBI\_TaxID=9940;

PROTEIN SEQUENCE.

RC TISSUE=Hypothalamus;  
 RX MEDLINE=9303624; PubMed=1417824;  
 RA Miyata A., Jiang L., Minamoto N., Arimura A.,  
 RT "Identification of calcitonin gene related peptide in ovine  
 hypothalamic extract.";  
 RL Biochem. Biophys. Res. Commun. 187:1474-1479(1992).  
 CC -! FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 neuromodulator role.  
 CC -! SUBCELLULAR LOCATION: Secreted.  
 CC -! SIMILARITY: Belongs to the calcitonin family.  
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 between the Swiss Institute of Bioinformatics and the EMBL outstation -  
 CC the European Bioinformatics Institute. There are no restrictions on its  
 CC use as long as its content is in no way modified and this statement is not  
 CC removed.

## PIR; JH0709; JH0709.

DR InterPro; IPR001693; Calcitonin-like.  
 DR Pfam; PF00214; Calc\_CGRP\_IAPP; I.  
 DR PRINTS; PR00817; CALCITONIN-B.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.  
 KW Amidation; Direct protein sequencing; Hormone.  
 FT MOD RES 37 37 Phenylalanine amide.  
 FT DISULFID 2 7 By similarity.  
 SQ SEQUENCE 37 AA; 3780 MW; F5DDF64D248B6A47 CRC64;

Query Match 42.1%; Score 85; DB 1; Length 37;  
 Best Local Similarity 50.0%; Pred. No. 0.00019;  
 Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

QY 2 CNTATCATQRIANFLVHSSNNFGPILPPTNVGSNTY 37  
 DB 2 CNTATCVTHRLAGLLSRSGGVKSNFVPTNVGSQAF 37

## RESULT 33

Q4THN9\_TETNG PRELIMINARY; PRT; 44 AA.  
 ID Q4THN9\_TETNG  
 AC Q4THN9;  
 DT 13-SEP-2005 (TREMBlrel. 31, Created)  
 DT 13-SEP-2005 (TREMBlrel. 31, Last sequence update)  
 DT 13-SEP-2005 (TREMBlrel. 31, Last annotation update)  
 DE Chromosome undetermined SCAF2800, whole genome shotgun sequence.  
 DE (Fragment).  
 GN ORFNames=GSTENG00000464001;  
 OS Tetraodon nigroviridis (Green puffer).

AC OAS173;  
 DT 13-SEP-2005 (TrEMBLrel. 31, Created)  
 DT 13-SEP-2005 (TrEMBLrel. 31, Last sequence update)  
 DT 13-SEP-2005 (TrEMBLrel. 31, Last annotation update)  
 DE Chromosome 5 SCAF14581, whole genome shotgun sequence.  
 DE (Fragment).  
 GN ORFNames=GSTENG00017812001;  
 OS Tetradon nigroviridis (Green puffer).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
 OC Acanthomorpha; Acanthopterygii; Percormorpha; Tetraodontiformes;  
 OC Tetraodontidae; Tetraodontidae; Tetraodon.  
 OC NCBI\_TaxID=99883;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RA Jaillon O., Aury J.M., Brunet F., Petit J.L., Strange-Thomann N.,  
 RA Micaud S., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,  
 RA Dastiva C., Salanoubat M., Levy M., Boudet N., Castellano S.,  
 RA Anthonard V., Jubin C., Castelli V., Katinka M., Vacherie B.,  
 RA Bleumont C., Skalli Z., Cactolico L., Poulin J., De Berardinis V.,  
 RA Parra G., Lardier G., Chapelle C., McKernan K.J., McMan P., Bosak S.,  
 RA Kellis M., Wolff J.N., Guigo R., Zody M.C., Mesirov J.,  
 RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,  
 RA Lauder V., Schachter V., Quelier F., Saurin W., Scarpelli C.,  
 RA Winkler P., Lander E.S., Weissenbach J., Roest Crolius H.;  
 RT "Genome duplication in the teleost fish Tetradon nigroviridis reveals  
 the early Vertebrate proto-karyotype.";  
 RL Nature 431:946-957(2004).  
 RU [2]  
 RP NUCLEOTIDE SEQUENCE.  
 RG Genoscope; Whitehead Institute Centre for Genome Research;  
 RL Submitted (FEB-2004) to the EMBL/GenBank/DBJ databases.  
 CC -1- CAUTION: The sequence shown here is derived from an  
 EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is  
 CC preliminary data.  
 DR EMBL; CA001014581; CAP99659.1; -; Genomic\_DNA.  
 FT NON\_TER 1 1  
 FT NON\_TER 51 51  
 SQ SEQUENCE 51 AA; 5551 MW; 45D08B150FC9FEE CRC64;  
 Query Match 43.6%; Score 88; DB 2; Length 51;  
 Best Local Similarity 50.0%; Pred. NO. 0.00011;  
 Matches 20; Conservative 4; Mismatches 8; Indels 8; Gaps 2;  
 QY 2 CMTATCACTORLANFLVHSS-----NNGPILPPTNVSNTY 37  
 DB 10 CMTATCVTHRLADPLSRSGSLGYSNF-----VPTNVAQAF 45  
 RESULT 30  
 CALCB HUMAN STANDARD; PRT; 127 AA.  
 ID P10092; O9UCN9;  
 AC P10092; O9UCN9;  
 DT 01-MAR-1989 (Rel. 10, Created)  
 DT 01-MAR-1989 (Rel. 10, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Calcitonin gene-related peptide II precursor (CGRP-II) (Beta-type  
 CGRP)  
 GN Name=CALCB; Synonyms=CALC2;  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homiidae;  
 OC Homo.  
 OC NCBI\_TaxID=9606;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=87105923; PubMed=3492393; DOI=10.1016/0014-5793(86)81091-2;  
 RA Steenbergh P.H., Hoepfener J.W.M., Zandberg J., Vissers A.,  
 RA Lips C.J.M., Jansz H.S.;  
 RT "Structure and expression of the human calcitonin/CGRP genes.";  
 RL FEBS Lett. 209:97-103(1986).

RN [2]  
 RP NUCLEOTIDE SEQUENCE OF 56-127.  
 RX MEDLINE=85180007; PubMed=2985435; DOI=10.1016/0014-5793(85)80820-6;  
 RA Steenbergh P.H., Hoepfener J.W.M., Zandberg J., Lips C.J.M.,  
 RA Jansz H.S.;  
 RT "A second human calcitonin/CGRP gene.";  
 RL FEBS Lett. 183:403-407(1985).  
 RN [3]  
 RP PARTIAL PROTEIN SEQUENCE OF 82-108.  
 RX MEDLINE=87109142; PubMed=3492492;  
 RA Petermann J.B., Born W., Chang J.Y., Fischer J.A.;  
 RT "Identification of a novel calcitonin gene-related peptide, and partial amino  
 acid sequence in the spinal cord.";  
 RL J. Biol. Chem. 262:542-545(1987).  
 RN [4]  
 RP PROTEIN SEQUENCE OF 82-86 AND 104-117.  
 RC TISSUE=Spinal cord;  
 RX MEDLINE=90211348; PubMed=2322288;  
 RA Wimalawansa S.J., Morris H.R., Etienne A., Blench I., Panico M.,  
 RA McIntyre I.;  
 RT "Isolation, purification and characterization of beta-hCGRP from human  
 spinal cord.";  
 RL Biochem. Biophys. Res. Commun. 167:993-1000(1990).  
 RN [5]  
 RP PROTEIN SEQUENCE OF 82-104.  
 RC TISSUE=Pheochromocytoma;  
 RX MEDLINE=92287083; PubMed=1318039;  
 RA Kitamura K., Kangawa K., Kawamoto M., Ichiki Y., Matsuo H., Eto T.;  
 RT "Isolation and characterization of peptides which act on rat  
 platelets, from a pheochromocytoma.";  
 RL Biochem. Biophys. Res. Commun. 185:134-141(1992).  
 CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 CC neuromodulator role.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
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 CC between the Swiss Institute of Bioinformatics and the EMBL outstation -  
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 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 DR EMBL; X04855; CAC05295.1; -; Genomic\_DNA.  
 DR EMBL; X04857; CAC05295.1; JOINED; Genomic\_DNA.  
 DR EMBL; X04861; CAC05295.1; JOINED; Genomic\_DNA.  
 DR EMBL; X02404; CAA26249.1; -; mRNA.  
 DR PIR; A25864; A25864.  
 DR PIR; I37232; I37232.  
 DR Ensembl; ENSG00000175868; Homo sapiens.  
 DR HGN; HGNC:1438; CALCB.  
 DR H-InvDB; HIX009469; -.  
 DR MIM; 114160; -.  
 DR GO; GO:0005625; C:soluble fraction; TAS.  
 DR GO; GO:0005184; P:neuropeptide hormone activity; TAS.  
 DR GO; GO:0006874; P:calcium ion homeostasis; TAS.  
 DR GO; GO:0007165; P:signal transduction; TAS.  
 DR InterPro; IPR001693; Calcitonin-B.  
 DR InterPro; IPR002163; Calcitonin-B.  
 DR Pfam; PF00214; Calc CGRP IAPP; I.  
 DR PRINTS; PR00817; CALCITONIN.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.  
 KW Amidation; Cleavage on pair of basic residues;  
 KW Direct protein sequencing; Hormone; Signal.  
 KW SIGNAL 25  
 FT PROPEP 1 26 79  
 FT PEPTIDE 82 118  
 FT PROPEP 124 127  
 FT MOD\_RES 118 118  
 FT Phenylalanine amide (G-119 provides amide group).

RP NUCLEOTIDE SEQUENCE.  
 RG Genoscope; Whitehead Institute Centre for Genome Research;  
 RL Submitted (FEB-2004) to the EMBL/GenBank/DBJ databases.  
 CC -! CAUTION: The sequence shown here is derived from an  
 CC EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is  
 CC preliminary data.  
 DR EMBL; CAAB01014769; CAG05615.1; -; Genomic\_DNA.  
 FT NON\_TER 184  
 SQ SEQUENCE 184 AA; 20107 MW; 851FB9A69FD16F29 CRC64;

Query Match 44.1%; Score 89; DB 2; Length 184;  
 Best Local Similarity 50.0%; Pred. No. 0.00024;  
 Matches 18; Conservative 3; Mismatches 15; Indels 0; Gaps 0;

QY 2 CNTATCATGRLANFLVHSSNNFGPIIPPTNVGSNTY 37  
 ||||| ||||| ||||| ||||| :  
 DB 140 CNTATCTVTHRLADFLRSQGMGNSNFPVPTNVGAKAF 175

RESULT 27  
 ID Q92164\_ONCSF PRELIMINARY; PRT; 56 AA.  
 AC Q92164;  
 DT 01-NOV-1996 (TREMBlrel. 01, Created)  
 DT 01-NOV-1996 (TREMBlrel. 01, Last sequence update)  
 DT 01-JUN-2003 (TREMBlrel. 24, Last annotation update)  
 DE Calcitonin gene-related peptide (Fragment).  
 GN Name=CGRP;  
 OS Oncorhynchus sp. (Salmon).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;  
 OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.  
 OX NCBI\_TaxID=8025;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=92344325; PubMed=1637123;  
 RA Janz H.S., Zandberg J.;  
 RT "Identification and partial characterization of the salmon  
 RL calcitonin/CGRP gene by polymerase chain reaction.",  
 RL Ann. N. Y. Acad. Sci. 657:63-69(1992).  
 DR EMBL; S40497; AAB22593.1; -; Genomic DNA.  
 DR GO; GO:0005576; C:extracellular region; IEA.  
 DR GO; GO:0005179; F:hormone activity; IEA.  
 DR InterPro; IPR001693; Calcitonin-like.  
 DR InterPro; IPR002163; Calcitonin B.  
 DR Pfam; PF00214; Calc\_CGRP\_IAPP; I.  
 DR PRINTS; PR00817; CALCITONINB.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.  
 FT NON\_TER 1  
 SQ SEQUENCE 56 AA; 6019 MW; C7852837BAF74314 CRC64;

Query Match 44.1%; Score 89; DB 2; Length 56;  
 Best Local Similarity 50.0%; Pred. No. 8.4e-05;  
 Matches 18; Conservative 3; Mismatches 15; Indels 0; Gaps 0;

QY 2 CNTATCATGRLANFLVHSSNNFGPIIPPTNVGSNTY 37  
 ||||| ||||| ||||| ||||| :  
 DB 8 CNTATCTVTHRLADFLRSQGMGNSNFPVPTNVGAKAF 43

RESULT 28  
 ID Q9M7Y1; STANDARD; PRT; 128 AA.  
 AC Q9M7Y1;  
 DT 28-FEB-2003 (Rel. 41, Created)  
 DT 28-FEB-2003 (Rel. 41, Last sequence update)  
 DT 10-MAY-2005 (Rel. 47, Last annotation update)  
 DE Calcitonin gene-related peptide I precursor (CGRP-1) (Alpha-type  
 DE CGRP).  
 GN Name=CALCA; Synonyms=CALC, CCALCI;  
 GN Canis familiaris (Dog).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

CC Mammalia; Euteria; Laurasiatheria; Carnivora; Fissipedia; Canidae;  
 CC Canis.  
 OX NCBI\_TaxID=9615;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=20424786; PubMed=10967131; DOI=10.1007/s003350010157;  
 RA Wende S., Krenn A., Breen M., Brunnberg L., Brenig B.;  
 RT "Molecular analysis and chromosomal assignment of the canine CALC-  
 RT I/alpha-CGRP gene.";  
 RL Mamm. Genome 11:736-740(2000).  
 RN [2]  
 RP PROTEIN SEQUENCE OF 26-50.  
 RC TISSUE=Thyroid;  
 RX MEDLINE=92100867; PubMed=1758974; DOI=10.1016/0167-0115(91)90082-R;  
 RA Mol J.A., Kwant M.M., Arnold I.C.J., Hazewinkel H.A.W.;  
 RT "Elucidation of the sequence of canine (pro)-calcitonin. A molecular  
 RT biological and protein chemical approach.";  
 RL Regul. Pept. 35:189-195(1991).  
 CC -! FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
 CC vessels including the coronary, cerebral and systemic vasculature.  
 CC Its abundance in the CNS also points toward a neurotransmitter or  
 CC neuromodulator role. It also elevates platelet cAMP (By  
 CC similarity).  
 CC -! SUBCELLULAR LOCATION: Secreted.  
 CC -! ALTERNATIVE PRODUCTS:  
 CC Event=Alternative splicing; Named isoforms=2;  
 CC Name=Calcitonin-gene related peptide I;  
 CC IsoId=Q9M7Y1-1; Sequence=Displayed;  
 CC Name=Calcitonin;  
 CC IsoId=P41547-1; Sequence=External;  
 CC -! SIMILARITY: Belongs to the calcitonin family.

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 CC removed.  
 CC -----  
 DR EMBL; AJ271090; CAB97487.1; -; Genomic DNA.  
 DR Ensembl; ENSCAF00000008562; Canis familiaris.  
 DR InterPro; IPR000443; Amy1in.  
 DR InterPro; IPR001693; Calcitonin-like.  
 DR InterPro; IPR002163; Calcitonin B.  
 DR Pfam; PF00214; Calc\_CGRP\_IAPP; I.  
 DR PRINTS; PR00817; CALCITONINB.  
 DR PRINTS; PR00818; ISLETAMYLOID.  
 DR SMART; SM00113; CALCITONIN; 1.  
 DR PROSITE; PS00258; CALCITONIN; 1.  
 KM Alternative splicing; Amidation; Cleavage on pair of basic residues;  
 KM Direct protein sequencing; Hormone; Signal.  
 FT SIGNAL 1 25  
 FT PROPEP 26 80 By similarity.  
 FT PEPTIDE 83 119 Calcitonin gene-related peptide I.  
 FT PROPEP 125 128 By similarity.  
 FT MOD\_RES 119 119 Phenylalanine amide (G-120 provides amide  
 FT DISULFID 84 89 group) (By similarity).  
 FT CONFLICT 45 45 E -> K (in Ref. 2).  
 FT CONFLICT 49 49 L -> N (in Ref. 2).  
 SQ SEQUENCE 128 AA; 13874 MW; 7233125BD8CB564 CRC64;

Query Match 44.1%; Score 89; DB 1; Length 128;  
 Best Local Similarity 52.5%; Pred. No. 0.00022;  
 Matches 21; Conservative 1; Mismatches 10; Indels 8; Gaps 2;

QY 2 CNTATCATGRLANFLVHSSNNFGPIIPPTNVGSNTY 37  
 ||||| ||||| ||||| ||||| :  
 DB 84 CNTATCTVTHRLADFLRSQGMGNSNFPVPTNVGAKAF 119

RESULT 29  
 ID Q4S173\_TETNG PRELIMINARY; PRT; 51 AA.

```

DR GO: GO:0005179; F: hormone activity; IEA.
DR InterPro: IPR000443; Amylin.
DR InterPro: IPR001693; Calcitonin-like.
DR Pfam: PF00214; Calc CGRP_IAPP; 1.
DR PRINTS: PR00818; ISLETAMYLOID.
DR SMART: SM00113; CALCITONIN; 1.
FT NON_TER 1 1
FT NON_TER 25 25
SQ SEQUENCE 25 AA; 2723 MW; 911B2C02PB677BAF CRC64;

Query Match 45.0%; Score 91; DB 2; Length 25;
Best Local Similarity 72.0%; Pred. No. 1.8e-05;
Matches 18; Conservative 2; Mismatches 5; Indels 0; Gaps 0;

QY 9 TORLANFLVHSSNFGPILPPTNVG 33
Db 1 TORLADFLVRSNNMGAFISPTNVG 25
|||||:|||||:|||||:|||||:
|T|L|A|D|F|L|V|R|S|N|N|M|G|A|F|I|S|P|T|N|V|G|

RESULT 24
ID P79814.ONCGO PRELIMINARY; PRT; 52 AA.
AC P79814;
DT 01-MAY-1997 (TrEMBLrel. 03, Created)
DT 01-MAY-1997 (TrEMBLrel. 03, Last sequence update)
DE 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE Calcitonin gene-related peptide 4 (Fragment).
OC Oncorhynchus gorbuscha (Pink salmon) (Humpback salmon).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;
OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
OX NCBI_TaxId=8017;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=97057244; Pubmed=6901583; DOI=10.1073/pnas.93.22.12344;
RA Janer H., Martal K., Zandberg U., Milhaud G., Benson A.A.,
RA Julienne A., Moukhtar M.S., Cressent M.;
RT "Identification of a new calcitonin gene in the salmon Oncorhynchus
RT gorbuscha."
RL Proc. Natl. Acad. Sci. U.S.A. 93:12344-12348(1996).
RN [2]
RP NUCLEOTIDE SEQUENCE.
RA Cressent M.D.;
RL Submitted (SEP-1996) to the EMBL/GenBank/DBJ databases.
DR EMBL: U71287; AAB38533.1; -; mRNA.
DR GO: GO:0005576; Extracellular region; IEA.
DR GO: GO:0005179; F: hormone activity; IEA.
DR InterPro: IPR001693; Calcitonin-like.
DR InterPro: IPR002163; Calcitonin_B.
DR Pfam: PF00214; Calc CGRP_IAPP; 1.
DR PRINTS: PR00817; CALCITONINB.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; 1.
FT CHAIN <1 4 N-terminal peptide.
FT CHAIN 7 43 calcitonin gene-related peptide 4.
FT CHAIN 49 52 carboxy terminal peptide.
FT NON_TER 1 1
SQ SEQUENCE 52 AA; 5700 MW; 3FAC471D2A682321 CRC64;

Query Match 44.6%; Score 90; DB 2; Length 52;
Best Local Similarity 50.0%; Pred. No. 5.6e-05;
Matches 18; Conservative 3; Mismatches 15; Indels 0; Gaps 0;

QY 2 CNTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37
Db 8 CNTATCVTRHLDPLRSRSGMGNSNFPPTNVGAKAF 43
|||||:|||||:|||||:|||||:
|C|N|T|A|T|C|A|T|O|R|A|N|F|L|V|H|S|S|N|F|G|P|I|L|P|P|T|N|V|G|S|N|T|Y|
|C|N|T|A|T|C|V|T|R|H|L|D|P|L|R|S|R|G|M|G|N|S|N|F|P|P|T|N|V|G|A|K|A|F|

RESULT 25
O80FT9_FUGRU PRELIMINARY; PRT; 126 AA.
ID O80FT9_FUGRU PRELIMINARY;
AC O80FT9;
DT 01-JUN-2002 (TrEMBLrel. 21, Created)

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DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE Calcitonin gene related peptide.
GN Name-cgfp;
OS Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;
OC Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
OC Tetraodontidae; Tetraodontidae; Takifugu.
OX NCBI_TaxId=31033;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Clark M.S.;
RL "Structure and expression of fugu calcitonin gene."
RL Submitted (MAR-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL: AJ309015; GAC81277.1; -; Genomic DNA.
DR Ensembl: SINEFRUG0000012598; Fugu rubripes.
DR GO: GO:0005576; Extracellular region; IEA.
DR GO: GO:0005179; F: hormone activity; IEA.
DR InterPro: IPR000443; Amylin.
DR InterPro: IPR001693; Calcitonin-like.
DR InterPro: IPR002163; Calcitonin_B.
DR Pfam: PF00214; Calc CGRP_IAPP; 1.
DR PRINTS: PR00817; CALCITONINB.
DR PRINTS: PR00818; ISLETAMYLOID.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; 1.
FT CHAIN 81 117 calcitonin gene related peptide.
SQ SEQUENCE 126 AA; 13863 MW; 31CB14A01B2CD57 CRC64;

Query Match 44.6%; Score 90; DB 2; Length 126;
Best Local Similarity 50.0%; Pred. No. 0.00015;
Matches 18; Conservative 3; Mismatches 15; Indels 0; Gaps 0;

QY 2 CNTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37
Db 82 CNTATCVTRHLDPLRSRSGMGNSNFPPTNVGAKAF 117
|||||:|||||:|||||:|||||:
|C|N|T|A|T|C|A|T|O|R|A|N|F|L|V|H|S|S|N|F|G|P|I|L|P|P|T|N|V|G|S|N|T|Y|
|C|N|T|A|T|C|V|T|R|H|L|D|P|L|R|S|R|G|M|G|N|S|N|F|P|P|T|N|V|G|A|K|A|F|

RESULT 26
O4S167_TERNG PRELIMINARY; PRT; 184 AA.
ID O4S167_TERNG PRELIMINARY;
AC O4S167;
DT 13-SEP-2005 (TrEMBLrel. 31, Created)
DT 13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT 13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE Chromosome 13 SCAP14769, whole genome shotgun sequence.
DE (Fragment).
GN ORFNames=GSTENG0025692001;
OS Tetraodon nigroviridis (Green puffer).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Neoteleostei;
OC Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
OC Tetraodontidae; Tetraodontidae; Tetraodon.
OX NCBI_TaxId=99883;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Jallion O., Aury J.M., Brunet F., Petit J.L., Strange-Thomann N.,
RA Mauceli E., Bouneau L., Fischer C., Ozouf-Coatac C., Bernot A.,
RA Nicada S., Jaffe D., Fleher S., Lutfalla G., Dossat C., Segurens B.,
RA Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano B.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Blemont C., Skalli Z., Catolico L., Poulain J., De Bernardis V.,
RA Cruaud C., Duprat S., Brottier P., Couranceau J.P., Gouzy J.,
RA Parra G., Lardier G., Chappie C., McKernan K.J., McEwan P., Bosak S.,
RA Kellis M., Volff J.N., Guigo R., Zody M.C., Mesirov J.,
RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA Laudet V., Schachter V., Quetier F., Saurin W., Scarpelli C.,
RA Winkler P., Lander E.S., Weissenbach J., Roest Crolius H.;
RT "Genome duplication in the teleost fish Tetraodon nigroviridis reveals
RT the early vertebrate proto-karyotype."
RN [2]

```

RP NUCLEOTIDE SEQUENCE, PROTEIN SEQUENCE OF 70-106, CHARACTERIZATION, AND  
RP MASS SPECTROMETRY.  
RC TISSUE=SKIN;  
RX MEDLINE=20148807; PubMed=10681586; DOI=10.1074/jbc.275.8.5934;  
RA Seon A.A., Pierre T.N., Redeker V., Lacombe C., Delfour A.,  
RA Nicolas P., Aniche M.,  
RT "Isolation, structure, synthesis, and activity of a new member of the  
RT calcitonin gene-related peptide family from frog skin and molecular  
RT cloning of its precursor."  
RL J. Biol. Chem. 275:5934-5940(2000).  
CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of  
CC vessels including the coronary, cerebral and systemic vasculature.  
CC Its abundance in the CNS also points toward a neurotransmitter or  
CC neuromodulator role (By similarity).  
CC -1- SUBCELLULAR LOCATION: Secreted.  
CC -1- TISSUE SPECIFICITY: Skin, intestine and brain.  
CC -1- MASS SPECTROMETRY: MW=3806.77; METHOD=MALDI; RANGE=70-106;  
CC NOTE=Ref.1.  
CC -1- SIMILARITY: Belongs to the calcitonin family.  
CC -----  
CC This Swiss-Prot entry is copyright. It is produced through a collaboration  
CC between the Swiss Institute of Bioinformatics and the EMBL outstation -  
CC the European Bioinformatics Institute. There are no restrictions on its  
CC use as long as its content is in no way modified and this statement is not  
CC removed.  
CC -----  
DR EMBL, Y18495; CAB76385.1; -; mRNA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR InterPro; IPR002163; Calcitonin B.  
DR Pfam; PF00214; Calc CGRP IAPP; 1.  
DR PRINTS; PR00817; CALCITONINB.  
DR PRINTS; PR00818; ISLETAMYLID.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; 1.  
KW Amidation; Cleavage on pair of basic residues;  
KW direct protein sequencing; Hormone; Signal.  
FT SIGNAL 1 25  
FT PROPEP 26 69  
FT CHAIN 70 106  
FT PROPEP 107 115  
FT MOD\_RES 106 106  
FT DISULFID 71 76  
FT SEQUENCE 115 AA; 12438 MW; A53D1125SCA53D31 CRC64;  
SQ  
Query Match 46.5%; Score 94; DB 1; Length 115;  
Best Local Similarity 50.0%; Pred. No. 3.8e-05;  
Matches 18; Conservative 7; Mismatches 11; Indels 0; Gaps 0;  
QY 2 CNTATCATGRLANFLVHSSNNFGPILPTNVGSNTY 37  
DB 71 CDTATCATGRLADFLRSRGIGSGKFPVPTVDSANF 106  
RESULT 22  
Q6DGJ9\_BRARE  
ID Q6DGJ9; BRARE PRELIMINARY; PRT; 126 AA.  
AC Q6DGJ9;  
DT 25-OCT-2004 (TREMBLrel. 28, Created)  
DT 25-OCT-2004 (TREMBLrel. 28, Last sequence update)  
DT 25-OCT-2004 (TREMBLrel. 28, Last annotation update)  
DE Zgc:92886.  
GN ORFName=zgc:92886;  
OS Brachydanio rerio (Zebrafish) (Danio rerio).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Cyprinidae; Danio.  
OX NCBI\_Taxid=7955;  
RN (1)  
RP NUCLEOTIDE SEQUENCE.

RC TISSUE=Brain;  
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.2426038999;  
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,  
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,  
RA Altschuler S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bat N.K.,  
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,  
RA Ditchenko L., Murnina K., Farmer A.A., Rubin G.M., Hong L.,  
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,  
RA Brownstein M.J., Ueda T.B., Toshynski S., Carninci P., Prange C.,  
RA Bask S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,  
RA Roach S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,  
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulik S.W.,  
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,  
RA Fahey J., Helton E., Kettelman M., Madan A., Rodriguez S., Sanchez A.,  
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,  
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,  
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,  
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smallos D.E.,  
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;  
RT "Generation and initial analysis of more than 15,000 full-length human  
RT and mouse cDNA sequences."  
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).  
RN [2]  
RP NUCLEOTIDE SEQUENCE.  
RC TISSUE=Brain;  
RA Strausberg R.;  
RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.  
DR EMBL; BC076343; AAH76343.1; -; mRNA.  
DR ZFIN; ZDB-GENE-040718-173; zgc:92886.  
DR GO; GO:0005576; C:extracellular region; IEA.  
DR GO; GO:0005179; P:hormone activity; IEA.  
DR InterPro; IPR000443; Amylin.  
DR InterPro; IPR001693; Calcitonin-like.  
DR InterPro; IPR002163; Calcitonin B.  
DR Pfam; PF00214; Calc CGRP IAPP; 1.  
DR PRINTS; PR00817; CALCITONINB.  
DR PRINTS; PR00818; ISLETAMYLID.  
DR SMART; SM00113; CALCITONIN; 1.  
DR PROSITE; PS00258; CALCITONIN; 1.  
DR SEQUENCE 126 AA; 13957 MW; 9A9399E3683D7B16 CRC64;  
SQ  
Query Match 45.5%; Score 92; DB 2; Length 126;  
Best Local Similarity 52.8%; Pred. No. 8e-05;  
Matches 19; Conservative 2; Mismatches 15; Indels 0; Gaps 0;  
QY 2 CNTATCATGRLANFLVHSSNNFGPILPTNVGSNTY 37  
DB 82 CNTATCVTHRLADFLRSRGIGSGKFPVPTVDSQAF 117  
RESULT 23  
Q9BBE1\_MACRU  
ID Q9BBE1; MACRU PRELIMINARY; PRT; 25 AA.  
AC Q9BBE1;  
DT 01-JUN-2001 (TREMBLrel. 17, Created)  
DT 01-JUN-2001 (TREMBLrel. 17, Last sequence update)  
DT 01-JUN-2003 (TREMBLrel. 24, Last annotation update)  
DE Islet amyloid polypeptide (Fragment).  
GN Name=iapp;  
OS Macropus rufus (Red kangaroo) (Megalala rufa).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Metatheria; Diprotodontia; Macropodidae; Macropus.  
OX NCBI\_Taxid=9521;  
RN (1)  
RP NUCLEOTIDE SEQUENCE.  
RA van Dijk M.A.M., de Jong W.W.;  
RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.  
RN [2]  
RP NUCLEOTIDE SEQUENCE.  
RA van Dijk M.A.;  
RL Submitted (FEB-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AJ268813; CAC28526.1; -; genomic DNA.  
DR GO; GO:0005576; C:extracellular region; IEA.

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RC TISSUE=Brain, and Intestine;
RX MEDLINE=93324452; PubMed=833553; DOI=10.1016/0196-9781(93)90148-A;
RA Conlon J.M., Tonon M.-C., Vaudry H.;
RT "Isolation and structural characterization of calcitonin gene-related
RL peptide from the brain and intestine of the frog, Rana ridibunda.";
RL PubMed 14:581-586(1993).
CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of
CC vessels including the coronary, cerebral and systemic vasculature.
CC its abundance in the CNS also points toward a neurotransmitter or
CC neuromodulator role.
CC -1- SIMILARITY: Belongs to the calcitonin family.
CC -----
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CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC the European Bioinformatics Institute. There are no restrictions on its
CC use as long as its content is in no way modified and this statement is not
CC removed.
CC -----
DR InterPro: IPR001693; Calcitonin-like.
DR InterPro: IPR002163; Calcitonin_B.
DR Pfam: PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS: PR00817; CALCITONINB.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; 1.
KW Amidation; Direct protein sequencing; Hormone.
FT MOD_RES 37 37 Phenylalanine amide.
FT DISUFID 2 7 By similarity.
SQ SEQUENCE 37 AA; 3887 MW; 0EEFE3AD2745EBDE CRC64;

Query Match 47.5%; Score 96; DB 1; Length 37;
Best Local Similarity 55.0%; Pred. No. 5.4e-06;
Matches 22; Conservative 2; Mismatches 8; Indels 8; Gaps 2;

QY 2 CNTATCATQRLANFLVHS---SNNFGPILPPTNGSNTY 37
Db 2 CNTATCVTHRLADFLRSQGVGNKF---VPTNGSKAF 37

RESULT 20
ID CALCA CHICK STANDARD; PRT; 125 AA.
AC P10286;
DT 01-MAR-1989 (Rel. 10, Created)
DT 01-NOV-1991 (Rel. 20, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Calcitonin gene-related peptide precursor (CGRP).
GN Name=CALCA; Synonyms=CALC;
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Archosauria; Aves; Neognathae; Galliformes; Phaethonidae; Phaethoninae;
OC Gallus.
OX NCBI_TaxID=9031;
RN (1)
RX MEDLINE=88030066; PubMed=3666142; DOI=10.1016/0014-5793(87)80510-0;
RA Minvielle S., Cressent M., Delehay M.C., Segond N., Milhaud G.,
RA Jullienne A., Mouttar M.S., Lasmole F.;
RT "Sequence and expression of the chicken calcitonin gene.";
RL FEBS Lett. 223:63-68(1987).
RN (2)
RP NUCLEOTIDE SEQUENCE OF 12-73.
RX MEDLINE=86030240; PubMed=4054101;
RA Lasmole F., Jullienne A., Day F., Minvielle S., Milhaud G.,
RA Mouttar M.S.;
RT "Cloning and the nucleotide sequence of chicken calcitonin mRNA:
RT direct evidence for the expression of a lower vertebrate calcitonin-
RL like gene in man and rat.";
RL EMBO J. 4:2603-2607(1985).
RN (3)
RP NUCLEOTIDE SEQUENCE OF 74-125.
RX MEDLINE=86248126; PubMed=3487468; DOI=10.1016/0014-5793(86)81425-9;
RA Minvielle S., Cressent M., Lasmole F., Jullienne A., Milhaud G.,
RA Mouttar M.S.;

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RT "Isolation and partial characterization of the calcitonin gene in a
RT lower vertebrate. Predicted structure of avian calcitonin gene-related
RL peptide.";
RL FEBS Lett. 203:7-10(1986).
CC -1- FUNCTION: CGRP induces vasodilatation. It dilates a variety of
CC vessels including the coronary, cerebral and systemic vasculature.
CC its abundance in the CNS also points toward a neurotransmitter or
CC neuromodulator role.
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- ALTERNATIVE PRODUCTS:
CC Event=Alternative splicing; Named isoforms=2;
CC Name=Calcitonin-gene related peptide;
CC IsoId=P10286-1; Sequence=Displayed;
CC Name=Calcitonin;
CC IsoId=P07650-1; Sequence=External;
CC -1- SIMILARITY: Belongs to the calcitonin family.
CC -----
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CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC the European Bioinformatics Institute. There are no restrictions on its
CC use as long as its content is in no way modified and this statement is not
CC removed.
CC -----
DR EMBL: X06311; CAA29630.1; -; Genomic_DNA.
DR EMBL: X06312; -; NOT_ANNOTATED_CDS; Genomic_DNA.
DR EMBL: X06314; CAA29633.1; -; Genomic_DNA.
DR EMBL: X03012; CAA26796.1; ALT TERM; mRNA.
DR EMBL: D00007; BAA00006.1; -; Genomic_DNA.
DR PIR: S00154; TCGRP.
DR EMBL: ENSGALG00000006054; Gallus gallus.
DR InterPro: IPR000443; Amylin.
DR InterPro: IPR001693; Calcitonin-like.
DR InterPro: IPR002163; Calcitonin_B.
DR Pfam: PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS: PR00817; CALCITONINB.
DR PRINTS: PR00818; ISLETAMYLTD.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; 1.
KW Alternative splicing; Amidation; Cleavage on pair of basic residues;
KW Hormone; Signal.
FT SIGNAL 1 25 Potential.
FT PROPEP 26 77 Calcitonin gene-related peptide.
FT PEPTIDE 80 116
FT PROPEP 122 125
FT MOD_RES 116 116 Phenylalanine amide (G-117 provides amide
FT group).
FT DISUFID 81 86 By similarity.
FT CONFLICT 56 56 D -> B (in Ref. 2).
SQ SEQUENCE 125 AA; 13729 MW; E864A2C9AC11F80 CRC64;

Query Match 47.5%; Score 96; DB 1; Length 125;
Best Local Similarity 55.0%; Pred. No. 2.2e-05;
Matches 22; Conservative 2; Mismatches 8; Indels 8; Gaps 2;

QY 2 CNTATCATQRLANFLVHS---SNNFGPILPPTNGSNTY 37
Db 81 CNTATCVTHRLADFLRSQGVGNKF---VPTNGSKAF 116

RESULT 21
ID CALCR_PHYBI STANDARD; PRT; 115 AA.
AC P81564;
DT 30-MAY-2000 (Rel. 39, Created)
DT 30-MAY-2000 (Rel. 39, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Skin calcitonin gene-related peptide precursor (S-CGRP).
OS Phylomedusa bicolor (Two-colored leaf frog).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Hylidae; Hylidae;
OC Phylomedusinae; Phylomedusa.
OX NCBI_TaxID=8393;
RN (1)

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SQ SEQUENCE 66 AA; 7131 MW; 3F0001BB7099770D CRC64;
Query Match 65.3%; Score 132; DB 2; Length 66;
Best Local Similarity 75.8%; Pred. No. 8.6e-11;
Matches 25; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVG 33
   :|||||:|||||:|||||:|||||:|||||:
DB 34 KCNTATCATQRLANFLVHSSNNFGALISPTDVG 66

RESULT 14
IAPP_SAGOE STANDARD; PRT; 32 AA.
AC Q28934;
DT 15-JUL-1998 (Rel. 36, Created)
DT 15-JUL-1998 (Rel. 36, Last sequence update)
DE 10-MAY-2005 (Rel. 47, Last annotation update)
DE Islet amyloid polypeptide (Amylin) (Fragment).
GN Name-IAPP;
OS Saginus oedipus (Cotton-top tamarin).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Primates; Platyrrhini;
OC Calitrichidae; Saginus.
OX NCBI_Taxid=9490;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Albrandt K., Sierrega M.E., Mull E., Brady E.M.G.;
RT "PCR amplification of amylin 3-34 from genomic DNA.";
RL Submitted (AUG-1996) to the EMBL/GenBank/DBJ databases.
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
CC -----
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CC use as long as its content is in no way modified and this statement is not
CC removed.
CC -----
CC EMBL; U62627; AAB05919.1; -; Genomic_DNA.
CC -----
CC DR InterPro; IPR000443; Amylin.
CC DR InterPro; IPR001693; Calcitonin-1like.
CC DR pfam; PF00214; Calc CGRP IAPP; 1.
CC DR PRINTS; PR00818; ISLETAMYL0ID.
CC DR SMART; SM00113; CALCITONIN; 1.
CC DR PROSITE; PS00258; CALCITONIN; PARTIAL.
CC DR Amyloid; Hormone.
CC KW Amyloid; Hormone.
CC FT PEPTIDE
CC FT NON_TER 1 1
CC FT NON_TER 32 32 Islet amyloid polypeptide.
CC FT NON_TER 32 32
CC SQ SEQUENCE 32 AA; 3340 MW; 91A219AEB3882C02 CRC64;

Query Match 59.4%; Score 120; DB 1; Length 32;
Best Local Similarity 75.0%; Pred. No. 1.9e-09;
Matches 24; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

QY 3 NTATCATQRLANFLVHSSNNFGPILPPTNVG 34
   |||||:|||||:|||||:|||||:|||||:
DB 1 NTATCSMRHLADFLGRSSNNFGALISPTNVG 32

RESULT 15
IAPP_PIG STANDARD; PRT; 32 AA.
AC Q29119;
DT 15-JUL-1998 (Rel. 36, Created)
DT 15-JUL-1998 (Rel. 36, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
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DE Islet amyloid polypeptide (Amylin) (Fragment).
GN Name-IAPP;
OS Sus scrofa (Pig).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suidae;
OC Sus.
OX NCBI_Taxid=9823;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Albrandt K., Sierrega M.E., Mull E., Brady E.M.G.;
RT "PCR amplification of amylin 3-34 from genomic DNA.";
RL Submitted (AUG-1996) to the EMBL/GenBank/DBJ databases.
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
CC -----
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CC removed.
CC -----
CC EMBL; U62628; AAB05919.1; -; Genomic_DNA.
CC -----
CC DR InterPro; IPR000443; Amylin.
CC DR InterPro; IPR001693; Calcitonin-1like.
CC DR pfam; PF00214; Calc CGRP IAPP; 1.
CC DR PRINTS; PR00818; ISLETAMYL0ID.
CC DR SMART; SM00113; CALCITONIN; 1.
CC DR PROSITE; PS00258; CALCITONIN; PARTIAL.
CC DR Amyloid; Hormone.
CC FT PEPTIDE
CC FT NON_TER 1 1
CC FT NON_TER 32 32 Islet amyloid polypeptide.
CC FT NON_TER 32 32
CC SQ SEQUENCE 32 AA; 3466 MW; 7EB37B990E555C8 CRC64;

Query Match 53.0%; Score 107; DB 1; Length 32;
Best Local Similarity 68.8%; Pred. No. 1.3e-07;
Matches 22; Conservative 0; Mismatches 10; Indels 0; Gaps 0;

QY 3 NTATCATQRLANFLVHSSNNFGPILPPTNVG 34
   |||||:|||||:|||||:|||||:|||||:
DB 1 NMATCATQRLANFLVHSSNNFGALISPTNVG 32

RESULT 16
IAPP_SHEEP STANDARD; PRT; 32 AA.
AC Q28605;
DT 15-JUL-1998 (Rel. 36, Created)
DT 15-JUL-1998 (Rel. 36, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Islet amyloid polypeptide (Amylin) (Fragment).
GN Name-IAPP;
OS Ovis aries (Sheep).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
OC Pecora; Bovidae; Caprinae; Ovis.
OX NCBI_Taxid=9940;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA Albrandt K., Sierrega M.E., Mull E., Brady E.M.G.;
RT "PCR amplification of amylin 3-34 from genomic DNA.";
RL Submitted (AUG-1996) to the EMBL/GenBank/DBJ databases.
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
CC -----
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```



```
RP NUCLEOTIDE SEQUENCE OF 1-66.
RA van Dijk M.A.M., de Jong W.W.;
RT "Indels indicate that rodents are monophyletic and lagomorphs are
   their sister group.";
RL Submitted (FEB-2000) to the EMBL/GenBank/DBJ databases.
RN [2]
RP NUCLEOTIDE SEQUENCE OF 36-67.
RA Albrandt K., Sierrega M.E., Mull E., Brady E.M.G.;
RT "PCR amplification of amylin 3-34 from genomic DNA.";
RL Submitted (Aug-1996) to the EMBL/GenBank/DBJ databases.
RN [3]
RP NUCLEOTIDE SEQUENCE OF 42-64.
RA STRAIN-New Zealand white;
RX MEDLINE=9315963; PubMed=8462765; DOI=10.1007/BF00399947;
RA Christensen L., Betscholtz C., Leckstroem A., Engstroem U., Corrie C.,
RA Johnson K.H., Adrian T.E., Westermarck P.;
RT "Islet amyloid polypeptide in the rabbit and European hare: studies on
   its relationship to amyloidogenesis.";
RL Diabetologia 36:183-188(1993).
CC -1- FUNCTION: Selectively inhibits insulin-stimulated glucose
   utilization and glycogen deposition in muscle, while not affecting
   adipocyte glucose metabolism.
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- SIMILARITY: Belongs to the calcitonin family.
CC -----
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   use as long as its content is in no way modified and this statement is not
   removed.
CC -----
DR EMBL: AJ286814; CAC28529.1; -; mRNA.
DR EMBL: U62630; AAB05917.1; -; Genomic_DNA.
DR EMBL: S57804; AAB26084.1; -; mRNA.
DR PIR: I46934; I46934.
DR InterPro: IPR000443; Amylin.
DR InterPro: IPR001693; Calcitonin-like.
DR Pfam: PF00214; Calc CGRP IAPP; 1.
DR PRINTS: PR00818; ISLETAMYL.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; PARTIAL.
KW Amyloid; Cleaveage on pair of basic residues; Hormone; Signal.
FT SIGNAL 1 22 Potential.
FT PROPEP 23 31
FT PEPTIDE 35 >67 Islet amyloid polypeptide.
FT DISULFD 35 40 By similarity.
FT NON_TER 67 67
SQ SEQUENCE 67 AA; 7230 MW; B5FEC2064F69646 CRC64;

Query Match 73.3%; Score 148; DB 1; Length 67;
Best Local Similarity 79.4%; Pred. No. 4,8e-13;
Matches 27; Conservative 2; Mismatches 5; Indels 0; Gaps 0;

Qy 1 KONTATCATQRLANFLVHSSNFGPILPPTVGS 34
Db 34 KNTVTTCATQRLANFLVHSSNFGAIFSPSPVGS 67

RESULT 12
QATB97_TETNG PRELIMINARY; PRT; 51 AA.
AC QATB97;
DT 13-SEP-2005 (TREMBLrel. 31, Created)
DT 13-SEP-2005 (TREMBLrel. 31, Last sequence update)
DE Chromosome undetermined SCAF7172, whole genome shotgun sequence.
DE (Fragment).
DE ORFNames=GSTENG0003849001;
OS Tetradon nigriviridis (Green puffer).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
OC Acanthomorpha; Acanthopterygii; Percomorpha; Tetradontiformes;
OC Tetradontidae; Tetradon.
```

```
OX NCBI_TaxID=99983;
RN NUCLEOTIDE SEQUENCE.
RA Tallon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,
RA Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
RA Nicoud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
RA Daellva C., Salanoubat M., Levy M., Boudet N., Castellano S.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Blomont C., Skalli Z., Catolico L., Poulain J., De Beraudins V.,
RA Craud C., Duprat S., Brotier P., Coutanceau J.P., Gouzy J.,
RA Parra G., Lardier G., Chapelle C., McKernan K.J., McEwan P., Bosak S.,
RA Kellis M., Volff J.N., Guigo R., Zody M.C., Mestrov J.,
RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA Lauder V., Schachter V., Quetier F., Saurin W., Scarpelli C.,
RA Winkler P., Lander E.S., Weissbach J., Roest Crolius H.;
RT "Genome duplication in the teleost fish Tetradon nigriviridis reveals
   the early vertebrate proto-karyotype.";
RL Nature 431:946-957(2004).
RN [2]
RP NUCLEOTIDE SEQUENCE.
RG GenomeScope; Whitehead Institute Centre for Genome Research;
RL Submitted (FEB-2004) to the EMBL/GenBank/DBJ databases.
CC -1- CAUTION: The sequence shown here is derived from an
   EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is
   preliminary data.
CC -----
DR EMBL: CA801007172; CAF89835.1; -; Genomic_DNA.
FT NON_TER 1 1
FT NON_TER 51 51
SQ SEQUENCE 51 AA; 5599 MW; 2AB836DCC4BBEF CRC64;

Query Match 70.8%; Score 143; DB 2; Length 51;
Best Local Similarity 73.0%; Pred. No. 1.8e-12;
Matches 27; Conservative 2; Mismatches 8; Indels 0; Gaps 0;

Qy 1 KONTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37
Db 8 KNTATCTQGLADFLVHSSNTIGTVAVPTVGSATY 44

RESULT 13
Q9BEFO_ERIEU PRELIMINARY; PRT; 66 AA.
AC Q9BEFO;
DT 01-JUN-2001 (TREMBLrel. 17, Created)
DT 01-JUN-2001 (TREMBLrel. 17, Last sequence update)
DT 01-MAR-2004 (TREMBLrel. 26, Last annotation update)
DE Islet amyloid polypeptide (Fragment).
DE Name=lapp;
GN Erinnaceus europaeus (Western European hedgehog).
OS Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Insectivora; Erinaceidae;
OC Erinaceinae; Erinaceus.
OX NCBI_TaxID=9365;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA van Dijk M.A.M., de Jong W.W.;
RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.
RN [2]
RP NUCLEOTIDE SEQUENCE.
RA van Dijk M.A.;
RL Submitted (FEB-2000) to the EMBL/GenBank/DBJ databases.
AC EMBL: AJ286815; CAC28520.1; -; mRNA.
DR GO: GO:0005576; C:extracellular region; IEA.
DR GO: GO:0005179; F:hormone activity; IEA.
DR InterPro: IPR000443; Amylin.
DR InterPro: IPR001693; Calcitonin-like.
DR InterPro: IPR002163; Calcitonin B.
DR Pfam: PF00214; Calc CGRP IAPP; 1.
DR PRINTS: PR00818; ISLETAMYL.
DR SMART: SM00113; CALCITONIN; 1.
DR PROSITE: PS00258; CALCITONIN; UNKNOWN_1.
FT NON_TER 66 66
```

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FT PROPE 77 91 Tyrosine amide (G-74 provides amide
FT MOD_RES 73 73 group).
FT DISULFID 38 43 By similarity.
SQ SEQUENCE 91 AA; 9925 MW; 42AB31AE1CE9EA99 CRC64;

Query Match
Best Local Similarity 81.1%; Score 160; DB 1; Length 91;
Matches 30; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTNGSNTY 37
Db 37 KCNTATCATQRLTNFLVRSNHLGALLPPTKVSNTY 73

RESULT 9
ID Q90743 CHICK PRELIMINARY; PRT; 135 AA.
AC Q90743
DT 01-NOV-1996 (TrEMBLrel. 01, Created)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE Amyloid protein precursor.
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC Gallus.
OX NCB1_TaxID=9031;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RC STRAIN=Leiborn; TISSUE=Pancrreas;
RX MEDLINE=95021303; PubMed=7935487; DOI=10.1210/me.8.6.713;
RA Fan L., Westermarck G., Chan S.J., Steiner D.F.;
RT "Altered gene structure and tissue expression of islet amyloid
RT polypeptide in the chicken.";
RL Mol. Endocrinol. 8:713-721(1994).
DR EMBL; L16955; AAA67704.1 -; mRNA.
DR PIR; A56855; A56855.
DR Ensemble; ENSGALG00000013168; Gallus gallus.
DR GO; GO:0005576; C:extracellular region; IEA.
DR GO; GO:0005524; F:ATP binding; IEA.
DR GO; GO:0005179; F:hormone activity; IEA.
DR GO; GO:0004672; F:protein kinase activity; IEA.
DR GO; GO:0006468; F:protein amino acid phosphorylation; IEA.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin-B.
DR InterPro; IPR000719; Prot kinase.
DR Pfam; PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS; PR00817; CALCITONIN.
DR PRINTS; PR00818; ISLETAMYLID.
DR ProDom; PD000001; Prot kinase; 1.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR Signal.
FT SIGNAL 1 22 Potential.
FT CHAIN 23 135 amyloid protein.
SQ SEQUENCE 135 AA; 14762 MW; 83DB3223AC735159 CRC64;

Query Match
Best Local Similarity 75.7%; Score 153; DB 2; Length 135;
Matches 29; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTNGSNTY 37
Db 80 KCNTATCATQRLADFLVRSNHLGALLPPTKVSNTY 116

RESULT 10
ID IAPP_CAVPO STANDARD; PRT; 92 AA.
AC IAPP_CAVPO
AC P12966;
DT 01-OCT-1989 (rel. 12, Created)

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DT 01-OCT-1989 (rel. 12, Last sequence update)
DT 10-MAY-2005 (rel. 47, Last annotation update)
DE Islet amyloid polypeptide precursor (Amylin).
GN Name=IAPP;
OS Cavia porcellus (Guinea pig).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
OC Hystriognathi; Caviidae; Cavia.
OX NCB1_TaxID=10141;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=89343542; PubMed=2668946;
RA Nishi M., Chan S.J., Nagamatsu S., Bell G.I., Steiner D.F.;
RT "Conservation of the sequence of islet amyloid polypeptide in five
RT mammals is consistent with its putative role as an islet hormone.";
RL Proc. Natl. Acad. Sci. U.S.A. 86:5738-5742(1989).
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
CC -----
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CC -----
DR EMBL; M25387; AAA7040.1 -; mRNA.
DR PIR; D3542; D3542.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR InterPro; IPR002163; Calcitonin-B.
DR Pfam; PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS; PR00817; CALCITONIN.
DR PRINTS; PR00818; ISLETAMYLID.
DR SMART; SM00113; CALCITONIN; 1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR AMIDATION; Amyloid; Cleavage on pair of basic residues; Hormone;
KW Signal.
FT SIGNAL 1 22 Potential.
FT PROPE 23 34 Islet amyloid polypeptide.
FT PEPTIDE 37 73
FT PROPE 77 92
FT MOD_RES 73 73 Tyrosine amide (G-74 provides amide
FT DISULFID 38 43 group) by similarity.
SQ SEQUENCE 92 AA; 9989 MW; 67F3629014BF3F9C CRC64;

Query Match
Best Local Similarity 74.8%; Score 151; DB 1; Length 92;
Matches 29; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTNGSNTY 37
Db 37 KCNTATCATQRLTNFLVRSNHLGALLPPTKVSNTY 73

RESULT 11
ID IAPP_RABIT STANDARD; PRT; 67 AA.
AC Q07334; Q28741; Q9BED7;
DT 15-JUL-1998 (rel. 36, Created)
DT 28-FEB-2003 (rel. 41, Last sequence update)
DT 10-MAY-2005 (rel. 47, Last annotation update)
DE Islet amyloid polypeptide precursor (Amylin) (Fragment).
GN Name=IAPP;
OS Oryctolagus cuniculus (Rabbit).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporidae;
OC Oryctolagus.
OX NCB1_TaxID=9986;
RN [1]

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DR PRINTS; PR00817; CALCITONINB.
DR SMART; SM00113; CALCITONIN.1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR AMADATION; Amyloid; Cleavage on pair of basic residues;
KW Direct protein sequencing; Hormone; Signal.
FT SIGNAL 1 22 Potential.
FT PROPEP 23 31 Islet amyloid polypeptide.
FT PEPTIDE 34 70
FT PROPEP 74 89
FT MOD_RES 70 70 Tyrosine amide (G-71 provides amide
FT DISULFID 35 40 group).
FT CONFLICT 67 67 By similarity.
SQ SEQUENCE 89 AA; 9832 MM; 0834D783DEAD72A8 CRC64;

Query Match 84.7%; Score 171; DB 1; Length 89;
Best Local Similarity 86.5%; Pred. No. 3.8e-16;
Matches 32; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37
DB 34 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 70

RESULT 7
ID IAPP_CANFA STANDARD; PRT; 89 AA.
AC P17716;
DT 01-AUG-1990 (Rel. 15, Created)
DT 01-FEB-1994 (Rel. 28, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Islet amyloid polypeptide precursor (Amylin).
GN Name=IAPP;
OS Canis familiaris (Dog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Carnivora; Fissipedia; Canidae;
OC Canis.
OC NCBI_Taxid=9615;
RX MEDLINE=92182022; PubMed=1543754; DOI=10.1016/0167-4781(92)90470-K;
RA Albrandt K., Mull E., Cooper G.J.S., Johnson M.J.;
RT "Nucleotide sequence of a cDNA for canine amylin."
RL Biochim. Biophys. Acta 1130:97-99(1992).
RN [2]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=90290487; PubMed=2192709;
RA Jordan K., Muttanah M.P., O'Brien T.D., Westermarck P., Betsholtz C.,
RA Johnson K.H.;
RT "Canine IAPP cDNA sequence provides important clues regarding
RT diabetogenesis and amyloidogenesis in type 2 diabetes."
RL Biochem. Biophys. Res. Commun. 169:502-508(1990).
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
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CC removed.
CC -----
CC EMBL; X59988; CAA42616.1; -; mRNA.
CC EMBL; M37720; AAA30849.1; -; mRNA.
CC PIR; S22344; S22344.
CC Ensemble; ENSCAG00000012365; Canis familiaris.
DR InterPro; IPR000443; Amylin.
DR InterPro; IPR001693; Calcitonin-like.
DR Pfam; PF00214; Calc_CGRP_IAPP; 1.
DR PRINTS; PR00817; CALCITONINB.
DR SMART; SM00113; CALCITONIN.1.
DR PROSITE; PS00258; CALCITONIN; 1.
KW Amyloid; Cleavage on pair of basic residues; Hormone;
FT SIGNAL 1 22 Potential.
FT PROPEP 23 34
FT PEPTIDE 37 73 Islet amyloid polypeptide.
```

```
DR PRINTS; PR00818; ISLETAMYLOID.
DR SMART; SM00113; CALCITONIN.1.
DR PROSITE; PS00258; CALCITONIN; 1.
DR AMADATION; Amyloid; Cleavage on pair of basic residues; Hormone;
KW Signal.
FT SIGNAL 1 22 Potential.
FT PROPEP 23 31 Islet amyloid polypeptide.
FT PEPTIDE 35 70
FT PROPEP 74 89
FT MOD_RES 70 70 Tyrosine amide (G-71 provides amide
FT DISULFID 35 40 group).
FT CONFLICT 67 67 By similarity.
SQ SEQUENCE 89 AA; 9800 MM; 9BF757E1C14935EF CRC64;

Query Match 83.7%; Score 169; DB 1; Length 89;
Best Local Similarity 86.5%; Pred. No. 7.2e-16;
Matches 32; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37
DB 34 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 70

RESULT 8
ID IAPP_OCTDE STANDARD; PRT; 91 AA.
AC P22889;
DT 01-AUG-1991 (Rel. 19, Created)
DT 01-AUG-1991 (Rel. 19, Last sequence update)
DT 10-MAY-2005 (Rel. 47, Last annotation update)
DE Islet amyloid polypeptide precursor (Amylin).
GN Name=IAPP;
OS Octodon degus (Degu).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
OC Hystricognathi; Octodontidae; Octodon.
OC NCBI_Taxid=10160;
RX MEDLINE=91155952; PubMed=2293024;
RA Nishi M., Steiner D.F.;
RT "Cloning of complementary DNAs encoding islet amyloid polypeptide,
RT insulin, and glucagon precursors from a New World rodent, the degu,
RT Octodon degus."
RL Mol. Endocrinol. 4:1192-1198(1990).
CC -!- FUNCTION: Selectively inhibits insulin-stimulated glucose
CC utilization and glycogen deposition in muscle, while not affecting
CC adipocyte glucose metabolism.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: Belongs to the calcitonin family.
CC This Swiss-Prot entry is copyright. It is produced through a collaboration
CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC the European Bioinformatics Institute. There are no restrictions on its
CC use as long as its content is in no way modified and this statement is not
CC removed.
CC -----
CC EMBL; M57669; AAA40589.1; -; mRNA.
CC PIR; A36118; A36118.
CC InterPro; IPR000443; Amylin.
CC InterPro; IPR001693; Calcitonin-like.
CC InterPro; IPR002163; Calcitonin B.
CC Pfam; PF00214; Calc_CGRP_IAPP; 1.
CC PRINTS; PR00817; CALCITONINB.
CC PRINTS; PR00818; ISLETAMYLOID.
CC SMART; SM00113; CALCITONIN.1.
CC PROSITE; PS00258; CALCITONIN; 1.
KW Amyloid; Cleavage on pair of basic residues; Hormone;
FT SIGNAL 1 22 Potential.
FT PROPEP 23 34
FT PEPTIDE 37 73 Islet amyloid polypeptide.
```

RP STRUCTURE BY NMR OF IAPP.  
 RX MEDLINE=91248117; PubMed=2039456;  
 RA Hubbard J.A.M., Martin S.R., Chaplin L.C., Bose C., Kelly S.M.,  
 RA Price N.C.;  
 RT "Solution structures of calcitonin-gene-related-peptide analogues of  
 RT calcitonin-gene-related peptide and amylin.";  
 RL Biochem. J. 275:785-788 (1991).  
 RN [12]  
 RP VARIANT GLY-53.  
 RX MEDLINE=96368727; PubMed=8772735;  
 RA Sakagashira S., Sanke T., Hanabusa T., Shimomura H., Ohagi S.,  
 RA Kumagaya K.Y., Nakajima K., Nango K.;  
 RT "Missense mutation of amylin gene (S20G) in Japanese NIDDM patients.";  
 RL Diabetes 45:1279-1281 (1996).  
 RN [13]  
 RP VARIANT GLY-53.  
 RX MEDLINE=99010531; PubMed=9794116; DOI=10.1007/s001250051060;  
 RA Chuang L.M., Lee K.C., Huang C.N., Wu H.P., Tai T.Y., Lin B.J.;  
 RT "Role of S20G mutation of amylin gene in insulin secretion, insulin  
 RT sensitivity, and type II diabetes mellitus in Taiwanese patients.";  
 RL Diabetologia 41:1250-1251 (1998).  
 CC -1- FUNCTION: Selectively inhibits insulin-stimulated glucose  
 CC utilization and glycogen deposition in muscle, while not affecting  
 CC adipocyte glucose metabolism.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- DISEASE: IAPP is the peptide subunit of amyloid found in  
 CC pancreatic islets of type 2 diabetic patients and in insulinomas.  
 CC -1- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
 CC This Swiss-Prot entry is copyright. It is produced through a collaboration  
 CC between the Swiss Institute of Bioinformatics and the EMBL outstation -  
 CC the European Bioinformatics Institute. There are no restrictions on its  
 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 DR EMBL, M27503; AAA35524.1; -; Genomic\_DNA.  
 DR EMBL, X14904; CAA33032.1; -; mRNA.  
 DR EMBL, X14905; CAA33033.1; -; mRNA.  
 DR EMBL, X14902; CAA33031.1; -; Genomic\_DNA.  
 DR EMBL, X14903; CAB57804.1; -; Genomic\_DNA.  
 DR EMBL, X13859; CAB57803.1; -; Genomic\_DNA.  
 DR EMBL, J04422; AAA52281.1; -; mRNA.  
 DR EMBL, M21785; AAA51728.1; -; Genomic\_DNA.  
 DR EMBL, M26550; AAA35983.1; -; Genomic\_DNA.  
 DR EMBL, X52818; CAA37002.1; -; Genomic\_DNA.  
 DR EMBL, X52819; CAA37002.1; JOINED; Genomic\_DNA.  
 DR EMBL, X56030; CAA39504.1; -; Genomic\_DNA.  
 DR EMBL, X55634; CAA39504.1; JOINED; Genomic\_DNA.  
 DR EMBL, X68830; CAA48724.1; -; Genomic\_DNA.  
 DR FIR, S04016; TCHUA.  
 DR Ensembl, ENSG00000121351; Homo sapiens.  
 DR HGNC, HGNC:5329; IAPP.  
 DR MIM, 147940; -;  
 DR GO, GO:0005625; C:soluble fraction; TAS.  
 DR GO, GO:0006102; F:receptor binding; TAS.  
 DR GO, GO:0006915; P:apoptosis; TAS.  
 DR GO, GO:0007267; P:cell-cell signaling; TAS.  
 DR GO, GO:0007165; P:signal transduction; TAS.  
 DR InterPro, IPR000443; Amylin.  
 DR InterPro, IPR001693; Calcitonin-like.  
 DR InterPro, IPR002163; Calcitonin-B.  
 DR Pfam, PF00214; Calc\_CGRP\_IAPP; 1.  
 DR PRINTS, PRO0817; CALCITONINB.  
 DR PRINTS, PRO0818; ISLETAMYLOID.  
 DR SMART, SM00113; CALCITONIN; 1.  
 DR PROSITE, PS00258; CALCITONIN; 1.  
 KW Amidation; Amyloid; Cleavage on pair of basic residues;  
 KW Direct protein sequencing; Hormone; Polymorphism; Signal.  
 FT SIGNAL 1 22  
 FT PROPEP 23 31  
 FT PEPTIDE 34 70  
 FT PROPEP 74 89  
 FT MOD\_RES 70 70  
 Tyrosine amide (G-71 provides amide

FT DISUPPID 35 40 (group).  
 FT VARIANT 53 53 S -> G (in dbSNP:1800203).  
 FT CONFLICT 53 53 /FTID=VAR\_012080.  
 FT SEQUENCE 89 AA; 9806 MW; AA8B1F7FD9FC4BD CRC64;  
 Query Match 88.1%; Score 178; DB 1; Length 89;  
 Best Local Similarity 91.9%; Pred. No. 3.9e-17;  
 Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
 QY 1 KCNTATCATGRLANFLVHSSNFGPILPPTVGSNTY 37  
 DB 34 KCNTATCATGRLANFLVHSSNFGALISSTVGSNTY 70  
 RESULT 6  
 IAPP\_FELCA  
 ID IAPP\_FELCA STANDARD; PRT; 89 AA.  
 AC P12967;  
 DT 01-OCT-1989 (Rel. 12, Created)  
 DT 01-OCT-1989 (Rel. 12, Last sequence update)  
 DT 13-SEP-2005 (Rel. 48, Last annotation update)  
 DE Islet amyloid polypeptide precursor (Amylin).  
 GN Name=IAPP;  
 OS Felis silvestris catus (Cat).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Laurasiatheria; Carnivora; Fissipedia; Felidae;  
 OC Felinae; Felis  
 OX NCBI\_TaxId=9685;  
 RN [1]  
 RP NUCLEOTIDE SEQUENCE.  
 RX MEDLINE=89345542; PubMed=2668946;  
 RA Nishi M., Chan S.J., Nagamatsu S., Bell G.I., Steiner D.F.;  
 RT "Conservation of the sequence of islet amyloid polypeptide in five  
 RT mammals is consistent with its putative role as an islet hormone.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 86:5738-5742 (1989).  
 RN [2]  
 RP PROTEIN SEQUENCE OF 34-50.  
 RX MEDLINE=87231921; PubMed=3035556;  
 RA Westermarck P., Westermarck C., Wlinder E., Hayden D.W., O'Brien T.D.,  
 RA Johnson K.H.;  
 RT "Amyloid fibrils in human insulinoma and islets of Langerhans of the  
 RT diabetic cat are derived from a neuropeptide-like protein also present  
 RT in normal islet cells.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 84:3881-3885 (1987).  
 RN [3]  
 RP NUCLEOTIDE SEQUENCE OF 34-70.  
 RX MEDLINE=91006862; PubMed=2210054;  
 RA Betsholtz C., Christmanon L., Engstrom U., Rozeman F., Jordan K.,  
 RA O'Brien T.D., Muraugh M., Johnson K.H., Westermarck P.;  
 RT "Structure of cat islet amyloid polypeptide and identification of  
 RT amino acid residues of potential significance for islet amyloid  
 RT formation.";  
 RL Diabetes 39:118-122 (1990).  
 CC -1- FUNCTION: Selectively inhibits insulin-stimulated glucose  
 CC utilization and glycogen deposition in muscle, while not affecting  
 CC adipocyte glucose metabolism.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the calcitonin family.  
 CC -----  
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 CC between the Swiss Institute of Bioinformatics and the EMBL outstation -  
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 CC use as long as its content is in no way modified and this statement is not  
 CC removed.  
 CC -----  
 DR EMBL, M25388; AAA30813.1; -; mRNA.  
 DR FIR, A33542; A33542.  
 DR InterPro, IPR000443; Amylin.  
 DR InterPro, IPR001693; Calcitonin-like.  
 DR InterPro, IPR002163; Calcitonin-B.  
 DR Pfam, PF00214; Calc\_CGRP\_IAPP; 1.

GenCore version 5.1.8  
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OM protein - protein search, using sw model

Run on: May 12, 2006, 15:14:16 ; Search time 186 Seconds  
(without alignments)  
87.403 Million cell updates/sec

Title: US-08-870-762b-1

Perfect score: 202

Sequence: 1 KCNTATCATQRLANFLVHSSNNFGPILPTNVGSNTY 37

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2443163 seqs, 439378781 residues

Total number of hits satisfying chosen parameters: 2443163

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 100 summaries

Database :

A\_Geneseq\_21:.\*  
1: geneseqp1980s:.\*  
2: geneseqp1990s:.\*  
3: geneseqp2000s:.\*  
4: geneseqp2001s:.\*  
5: geneseqp2002s:.\*  
6: geneseqp2003as:.\*  
7: geneseqp2003bs:.\*  
8: geneseqp2004s:.\*  
9: geneseqp2005s:.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	202	100.0	37	2	AAR29199 Pro(25), 28
2	202	100.0	37	2	AAR37792 Pro(25), Pr
3	202	100.0	37	2	AAM22540 Amylin ag
4	202	100.0	37	2	AAM74607 Amylin ag
5	202	100.0	37	2	AAM90165 Human amy
6	202	100.0	37	2	AAV22437 Pro(25), Pr
7	202	100.0	37	2	AAW90144 Human amy
8	202	100.0	37	3	AAB18572 Amino aci
9	202	100.0	37	5	ABR05489 Human amy
10	202	100.0	37	7	ADP17060 Human alb
11	202	100.0	37	8	AD136171 Human amy
12	202	100.0	37	8	ADN16159 Human pep
13	202	100.0	37	8	ADO51015 Human amy
14	202	100.0	37	9	ADV92867 Human amy
15	202	100.0	37	9	ADV92870 Amylin pe
16	202	100.0	37	9	AEI17936 Human amy
17	202	100.0	37	9	AEI17969 Human amy
18	201	99.5	37	2	AAAR29213 Pro(25) Va
19	201	99.5	37	2	AAAR29212 Ile(17) Pr
20	201	99.5	37	2	AAAR38817 Pro(25), Va
21	201	99.5	37	2	AAAR38816 Ile(17), Pr
22	201	99.5	37	2	AAV22439 Pro(25), Va
23	201	99.5	37	2	AAV22449 Ile(17), Pr
24	201	99.5	37	2	AAW90146 Human amy

25	201	99.5	37	2	AAW90154 Human amy
26	201	99.5	37	3	AAB18578 Amino aci
27	201	99.5	37	3	AAB18589 Amino aci
28	201	99.5	37	5	ABB05492 Human amy
29	201	99.5	37	5	ABB05504 Human amy
30	201	99.5	37	8	AD136177 Human amy
31	201	99.5	37	8	AD136188 Human amy
32	201	99.5	37	8	AD051032 Human amy
33	201	99.5	37	8	AD051021 Human amy
34	201	99.5	37	9	AEI17939 Human amy
35	201	99.5	37	9	AEI17969 Human amy
36	199	98.5	37	2	AAW22541 Amylin ag
37	199	98.5	36	2	AAAR29204 Des-Lys(1
38	197	97.5	36	2	AAAR37793 Des-Lys1
39	197	97.5	36	2	AAV22442 Des-Lys1,
40	197	97.5	36	2	AAW90145 Human amy
41	197	97.5	36	3	AAB18581 Amino aci
42	197	97.5	36	5	ABB05495 Human amy
43	197	97.5	36	8	AD136180 Human amy
44	197	97.5	36	8	AD051024 Human amy
45	196	97.0	36	2	AAAR37794 Des-Lys1
46	196	97.0	37	2	AAW22579 Amylin ag
47	195	96.5	36	7	ADH22050 Amylin an
48	195	96.5	37	2	AAAR29205 Leu(23) Pr
49	195	96.5	37	2	AAAR29211 Ile(17) Le
50	195	96.5	37	2	AAAR38809 Leu(23), Pr
51	195	96.5	37	2	AAAR38815 Ile17, Le
52	195	96.5	37	2	AAV22448 Ile17, Le
53	195	96.5	37	2	AAW90153 Human amy
54	195	96.5	37	2	AAW90147 Human amy
55	195	96.5	37	3	AAB18582 Amino aci
56	195	96.5	37	3	AAB18588 Amino aci
57	195	96.5	37	5	ABB05497 Human amy
58	195	96.5	37	5	ABB05503 Human amy
59	195	96.5	37	8	AD136187 Human amy
60	195	96.5	37	8	AD136181 Human amy
61	195	96.5	37	8	AD051025 Human amy
62	195	96.5	37	8	AD051031 Human amy
63	195	96.5	37	9	ADV92850 Human amy
64	195	96.5	37	9	ADV92844 Amylin pe
65	195	96.5	37	9	AEI17943 Human amy
66	195	96.5	37	9	AEI17949 Human amy
67	194	96.0	37	2	AAAR37786 Pro(25), Pr
68	194	96.0	37	2	AAAR37790 Pro(25), Pr
69	194	96.0	37	2	AAV22440 Arg18, Pr
70	194	96.0	37	2	AAW90142 Human amy
71	194	96.0	37	3	AAB18579 Amino aci
72	194	96.0	37	5	ABB05493 Human amy
73	194	96.0	37	8	AD136178 Human amy
74	194	96.0	37	8	AD051022 Human amy
75	194	96.0	37	9	ADV92841 Amylin pe
76	194	96.0	37	9	ADV92866 Amylin pe
77	194	96.0	37	9	AEI17940 Human amy
78	194	96.0	37	9	AEI17965 Human amy
79	193	95.5	36	9	ADV92843 Amylin pe
80	193	95.5	36	9	AEI17942 Human amy
81	193	95.5	37	2	AAAR29203 Des-Lys(1
82	193	95.5	37	2	AAW22551 Amylin ag
83	193	95.5	37	5	ABB05496 Human amy
84	192	95.0	36	9	ADV92868 Amylin pe
85	192	95.0	36	9	AEI17967 Human amy
86	192	95.0	37	2	AAW22577 Amylin ag
87	191.5	94.8	38	9	AEI17970 Human amy
88	191	94.6	37	9	AEI17950 Human amy
89	191	94.6	37	9	AEI17969 Human amy
90	190	94.1	36	2	AAAR29213 Des-Lys(1
91	190	94.1	36	2	AAAR38817 Des-Lys1
92	190	94.1	36	2	AAV22450 Des-Lys1,
93	190	94.1	36	2	AAW90155 Human amy
94	190	94.1	36	3	AAB18590 Amino aci
95	190	94.1	36	5	ABB05505 Human amy
96	190	94.1	36	8	AD136189 Human amy
97	190	94.1	36	8	AD051033 Human amy

98 190 94.1 36 9 ADV92852  
99 190 94.1 36 9 AEB17951  
100 189 93.6 36 2 AAR37791

Adv92852 Amylin pe  
Aeb17951 Human amy  
Aar37791 (Des-Lys1

## ALIGNMENTS

## RESULT 1

AAR29199  
ID AAR29199 standard; protein; 37 AA.

XX AAR29199;

XX 25-MAR-2003 (revised)

DT 20-APR-1993 (first entry)

XX Pro(25,28,29)-h-amylin for treating anorexia.

XX Anorexia; cachexia; adipose; amylin.

XX Homo sapiens.

XX WO9220367-A1.

XX 26-NOV-1992.

XX 23-MAY-1992; 92WO-US004357.

XX 24-MAY-1991; 91US-00704995.

XX 03-APR-1992; 92US-00862500.

XX (AMYL-) AMYLIN PHARM INC.

XX Rink TJ, Young AA;

XX WPI; 1992-415470/50.

XX Use of amylin and opt. insulin for treating anorexia - increases plasma

XX amylin and/or insulin levels, also for treating cachexia conditions,

XX adipose tissue deficiency etc.

XX Disclosure; Page 16; 19pp; English.

XX Treatment of a patient with anorexia or related condition comprises

XX admin. of amylin or an analogue in an amt. sufficient to increase the

XX amylin level in the plasma of the patient. The pref. amylin analogues are

XX given in AAR29197-222. Treating a patient deficient in adipose tissue

XX comprises admin. of amylin or an analogue and/or insulin in an amt. and

XX ratio sufficient to increase adipose tissue. Typical dosage units contain

XX 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-

XX 2003 to correct PN field.)

XX SQ Sequence 37 AA;

## RESULT 2

AAR37792  
ID AAR37792 standard; peptide; 37 AA.

XX AAR37792;

XX 25-MAR-2003 (revised)

DT 07-SEP-1993 (first entry)

XX

DE Pro25, Pro28, Pro29 human amylin analogue.  
XX Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;  
XX hyperglycaemic agent.

XX Homo sapiens.

XX Key Location/Qualifiers

XX Disulfide-bond 2..7

XX Misc-difference 25 /note= "Pro replaces wild-type Ala"

XX FT Misc-difference 28 /note= "Pro replaces wild-type Ser"

XX FT Misc-difference 29 /note= "Pro replaces wild-type Ser"

XX FT Modified-site 37 /note= "amidated"

XX WO9310146-A1.

XX 27-MAY-1993.

XX 19-NOV-1992; 92WO-US009842.

XX 19-NOV-1991; 91US-00794266.

XX (AMYL-) AMYLIN PHARM INC.

XX Gaeta LSL, Jones H, Albrecht E;

XX WPI; 1993-182488/22.

XX New amylin agonist peptide(s) - used for treatment and prevention of

XX hypoglycaemia and diabetes mellitus.

XX Claim 43; Fig 1 and Page 22; 43pp; English.

XX This peptide is an example of amylin agonists of the invention which can

XX be used as hyperglycaemics. The peptide is an analogue of human amylin

XX which mimics the effects of the wild-type hormone. Preferred peptides are

XX used in admixture with insulin for the treatment of diabetes mellitus or

XX with glucagon for the treatment of hypoglycaemic conditions. See AAR37799

XX CC CC and AAR38809-R38826. (Updated on 25-MAR-2003 to correct PN

XX field.)

XX SQ Sequence 37 AA;

XX Query Match 100.0%; Score 202; DB 2; Length 37;

XX Best Local Similarity 100.0%; Pred. No. 6.4e-21;

XX Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

XX 1 KONTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37

XX 1 KONTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37

XX RESULT 3

XX AAM22540

XX AAM22540 standard; peptide; 37 AA.

XX 12-OCT-1997 (first entry)

XX Amylin agonist peptide 25,28,29-Pro-h-amylin.

XX appetite regulation; amylin agonist; hybrid; CCK; cholecystokinin.

XX Synthetic.

XX Key Location/Qualifiers

XX Disulfide-bond 2..7

XX Modified-site 37



Query Match 100.0%; Score 202; DB 2; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37

RESULT 6  
AAV22437

ID AAV22437 standard; peptide; 37 AA.

AC AAV22437;

DT 28-SEP-1999 (first entry)

DE Pro25, Pro28, Pro29 amylin analogue.

KW Amylin agonist; human; insulin; diabetes; post-prandial glucose level;  
therapy; mutein.

OS Homo sapiens.

OS Synthetic.

Key Location/Qualifiers

FT Disulfide-bond 2..7

FT Misc-difference 25

FT Misc-difference 28

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

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FT Misc-difference 29

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FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37

RESULT 7  
AAW90144

ID AAW90144 standard; peptide; 37 AA.

AC AAW90144;

DT 15-MAR-1999 (first entry)

DE Human amylin agonist peptide 25,28,29-Pro-amylin.

KW Amylin; human; agonist; gastritis; gastric ulceration; treatment;

KW non-steroidal anti-inflammatory agent; NSAID; intravenous; subcutaneous;

KW pain; fever; inflammation; arthritis; hypercoagulation.

OS Homo sapiens.

OS Synthetic.

Key Location/Qualifiers

FT Disulfide-bond 2..7

FT Misc-difference 25

FT Misc-difference 28

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

FT Misc-difference 29

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FT Misc-difference 29

RESULT 8

Query Match 100.0%; Score 202; DB 2; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37



AB18572  
ID AAB18572 standard; peptide; 37 AA.  
XX  
AC AAB18572;  
XX  
DT 15-JAN-2001 (first entry)  
XX  
DE Amino acid sequence of an amylin agonist analogue compound.  
XX  
KM Amylin agonist; amylin; gastric motility; gastric emptying;  
KM postprandial dumping syndrome; postprandial hyperglycemia;  
KM gastrointestinal diagnostic procedure; gastrointestinal disorder; spasm;  
KM acute diverticulitis; biliary tract disorder; sphincter of oddi disorder.  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Disulfide-bond 2..7  
FT Modified-site 37  
FT /note="amidated residue"  
XX  
PN US6114304-A.  
XX  
PD 05-SEP-2000.  
XX  
PF 07-SEP-1994; 94US-00302069.  
XX  
PR 07-SEP-1993; 93US-00118381.  
XX  
PA (AMYL-) AMYLIN PHARM INC.  
XX  
PI Young AA, Rink TJ, Brown KAK, Koltzman OG;  
XX  
DR WPI; 2000-601336/57.  
XX  
PT Treating gastrointestinal disorder e.g. spasm by reducing gastric  
PT motility or delaying gastric emptying, postprandial dumping syndrome or  
PT postprandial hyperglycemia, by administering amylin or amylin agonist.  
XX  
PS Disclosure; Col 35-36; 50pp; English.  
XX  
XX The present sequence represents an amylin agonist analogue compound.  
CC Amylin or amylin agonists are administered for reducing gastric motility  
CC or delaying gastric emptying, and for treating postprandial dumping  
CC syndrome or postprandial hyperglycemia, by inducing amylin activity, in a  
CC mammal. The peptides are used to reduce gastric motility or for delaying  
CC gastric emptying in a mammal undergoing gastrointestinal diagnostic  
CC procedures, such as radiological examination or magnetic resonance  
CC imaging. They are also used for reducing gastric motility in  
CC gastrointestinal disorder, especially spasm, which is associated with a  
CC disorder of acute diverticulitis or disorders of biliary tract or  
CC sphincter of oddi. They are also used to treat postprandial dumping  
CC syndrome or postprandial hyperglycemia  
XX  
SQ Sequence 37 AA;

Query Match 100.0%; Score 202; DB 3; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 9  
ID ABB05489  
XX ABB05489 standard; peptide; 37 AA.  
XX  
AC ABB05489;  
XX  
DT 19-APR-2002 (first entry)  
XX

DE Human amylin agonist 25,28,29Pro-h-amylin, pramlintide.  
XX  
XX Human; amylin agonist; amylin peptide analogue; acetate; carbohydrate;  
KM liquid pharmaceutical formulation; polyhydric alcohol; phosphate;  
KM citrate; glutamate; buffer; antidiabetic; type II diabetes.  
XX  
OS Homo sapiens.  
OS Synthetic.  
XX  
PN US2001043934-A1.  
XX  
PD 22-NOV-2001.  
XX  
PF 09-JAN-1998; 98US-00005262.  
XX  
PR 08-JAN-1997; 97US-0035140P.  
XX  
PA (LITA/) L'ITALIEN J.  
PA (MUSU/) MUSUNURI S.  
PA (RUBY/) RUBY K.  
XX  
PI L'italien J, Musunuri S, Ruby K;  
XX  
DR WPI; 2002-163554/21.  
XX  
PT New pharmaceutical formulation useful for treating patients with type II  
PT diabetes comprises amylin agonist, carbohydrate or polyhydric alcohol and  
PT a buffer.  
XX  
PS Example 1; Page; 19pp; English.  
XX  
CC The present invention describes a liquid pharmaceutical formulation (A)  
CC comprising (wt/vol%) an amylin agonist (0.01 - 0.5), a carbohydrate or a  
CC polyhydric alcohol (1 - 10) and an acetate, phosphate, citrate or  
CC glutamate buffer (0.02 - 0.5) having a pH of 3 - 6. Also described is a  
CC commercial package containing the liquid pharmaceutical formulation (A).  
CC The package comprises a borosilicate glass vial having an open end, a  
CC stopper for multise compatible with the amylin and/or amylin agonist  
CC fixed in the open end of the vial and an aluminum band to retain the  
CC stopper in the far end of the vial. The package also comprises a  
CC cartridge for use in a pen injector. (A) has antidiabetic activity and  
CC can be used in the treatment of patients with type II diabetes. The  
CC formulation comprises amylin agonist which is biologically active, has a  
CC reduced tendency to form aggregates in water or at a pressure of greater  
CC than 2 psi and has a reduced tendency to precipitate in the presence of  
CC NaCl compared to human amylin. The formulation maintains stability upon  
CC storage under refrigerated or room-temperature conditions. The  
CC formulation retains short-term mixing compatibility with insulin and  
CC results in improved stability of the hormone and the patients no longer  
CC need to refrigerate the vial of insulin in use. The present sequence  
CC represents a human amylin peptide analogue (called pramlintide), which  
CC can be used as an amylin agonist in the present invention. N.B. The  
CC present sequence is not given in the present specification but is derived  
CC from the 37 amino acid human amylin as stated in the invention  
XX  
SQ Sequence 37 AA;

Query Match 100.0%; Score 202; DB 5; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 10  
ID ADF17060  
XX ADF17060 standard; protein; 37 AA.  
XX  
AC ADF17060;  
XX  
DT 12-FEB-2004 (first entry)  
XX

XX DE Human albumin fusion protein-related protein SegID2200.  
 XX DE albumin fusion protein; albumin activity; human serum albumin;  
 KW serum osmotic pressure; shelf-life; stability; antidiabetic;  
 KW gene therapy; diabetes mellitus; human.  
 XX  
 OS Homo sapiens.  
 XX WO2003060071-A2.  
 XX PD 24-JUL-2003.  
 XX PF 23-DEC-2002; 2002WO-US040891.  
 XX PR 21-DEC-2001; 2001US-034181P.  
 XX PR 24-JAN-2002; 2002US-0350358P.  
 XX PR 28-JAN-2002; 2002US-0351360P.  
 XX PR 26-FEB-2002; 2002US-0359370P.  
 XX PR 28-FEB-2002; 2002US-036000P.  
 XX PR 27-MAR-2002; 2002US-037500P.  
 XX PR 08-APR-2002; 2002US-0370227P.  
 XX PR 10-MAY-2002; 2002US-0378950P.  
 XX PR 24-MAY-2002; 2002US-0382617P.  
 XX PR 28-MAY-2002; 2002US-0383123P.  
 XX PR 05-JUN-2002; 2002US-0385708P.  
 XX PR 10-JUL-2002; 2002US-0394625P.  
 XX PR 24-JUL-2002; 2002US-0398008P.  
 XX PR 09-AUG-2002; 2002US-0402131P.  
 XX PR 13-AUG-2002; 2002US-0402708P.  
 XX PR 18-SEP-2002; 2002US-0411355P.  
 XX PR 18-SEP-2002; 2002US-0411426P.  
 XX PR 02-OCT-2002; 2002US-0414984P.  
 XX PR 11-OCT-2002; 2002US-0417611P.  
 XX PR 23-OCT-2002; 2002US-0420246P.  
 XX PR 05-NOV-2002; 2002US-0423623P.  
 XX PA (HUMA-) HUMAN GENOME SCI INC.  
 XX PA (DEL2) DELTA BIOTECHNOLOGY LTD.  
 XX PA (PRIN-) PRINCIPIA PHARM CORP.  
 XX PI Balance DJ, Turner AJ, Rosen CA, Haseltine WA;  
 XX DR WPI; 2003-598517/56.  
 XX PR New albumin fusion protein, useful for preparing a composition for  
 XX PT treating diabetes mellitus.  
 XX PS Example 4; SEQ ID NO 2200; 24pp; English.  
 XX CC This invention relates to a novel albumin fusion protein having albumin  
 XX CC or biological activity. Human serum albumin is responsible for a  
 XX CC significant proportion of the osmotic pressure of serum and also  
 XX CC functions as a carrier of endogenous and exogenous ligands. The fusion of  
 XX CC albumin to a therapeutic protein may increase shelf-life and stability of  
 XX CC the therapeutic protein. The albumin fusion protein of the invention may  
 XX CC allow production of compositions with antidiabetic activity whilst the  
 XX CC nucleotide sequence which encodes it may be useful for gene therapy. The  
 XX CC albumin fusion protein is useful for preparing a composition for treating  
 XX CC diabetes mellitus. The present sequence is that of a therapeutic protein  
 XX CC which was fused with human albumin to create a novel albumin fusion  
 XX CC protein of the invention. Note: The sequence data for this patent did not  
 XX CC form part of the printed specification, but was obtained in electronic  
 XX CC format directly from WIPO at ftp.wipo.int/pub/publishedpct\_sequences  
 XX SQ Sequence 37 AA;  
 QY Query Match 100.0%; Score 202; DB 7; Length 37;  
 Best Local Similarity 100.0%; Pred. No. 6,4e-21;  
 Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 1 KCNTATCATORLANFLVHSSNNFGPILPTTVGSGNTY 37  
 ||||||||||||||||||||||||||||||||||||||||

DB 1 KCNTATCATORLANFLVHSSNNFGPILPTTVGSGNTY 37  
 RESULT 11  
 AD136171  
 ID AD136171 standard; peptide; 37 AA.  
 XX AC AD136171;  
 XX DT 15-APR-2004 (first entry)  
 XX DE Human amylin agonist analogue #1.  
 XX KW Gastric motility; gastric emptying; amylin agonist; therapy; diagnosis;  
 KW antidiabetic; hypoglycaemia; human.  
 XX OS Homo sapiens.  
 XX FH Key Location/Qualifiers  
 FT Disulfide-bond 2..7  
 FT Modified-site 37  
 FT /note= "Amidated tyrosine"  
 XX PN US6608029-B1.  
 XX PD 19-AUG-2003.  
 XX PR 22-MAY-2000; 2000US-00576062.  
 XX PR 07-SEP-1993; 93US-00118361.  
 XX PR 07-SEP-1994; 94US-00302069.  
 XX PA (AMYL-) AMYLIN PHARM INC.  
 XX PI Kolterman OG, Young AA, Rink TJ, Keating Brown KA;  
 XX DR WPI; 2004-118064/12.  
 XX PT Reducing gastric motility or delaying gastric emptying in a mammal,  
 PT useful for treating post-prandial hyperglycemia, comprises administering  
 PT an amylin or an amylin agonist.  
 XX PS Disclosure; SEQ ID NO 1; 51pp; English.  
 XX CC The present invention is directed to novel methods for reducing gastric  
 XX CC motility and delaying gastric emptying, comprising the administration of  
 XX CC an amylin or an amylin agonist. The invention is useful for reducing  
 XX CC gastric motility and delaying gastric emptying for therapeutic and  
 XX CC diagnostic purposes. The invention is also useful for treating conditions  
 XX CC associated with elevated, inappropriate or undesired post-prandial blood  
 XX CC glucose levels and treating ingestion of a toxin. The present sequence is  
 XX CC human amylin agonist analogue.  
 XX SQ Sequence 37 AA;  
 QY Query Match 100.0%; Score 202; DB 8; Length 37;  
 Best Local Similarity 100.0%; Pred. No. 6,4e-21;  
 Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 1 KCNTATCATORLANFLVHSSNNFGPILPTTVGSGNTY 37  
 ||||||||||||||||||||||||||||||||||||||||  
 DB 1 KCNTATCATORLANFLVHSSNNFGPILPTTVGSGNTY 37  
 ||||||||||||||||||||||||||||||||||||||||  
 RESULT 12  
 ADN16159  
 ID ADN16159 standard; peptide; 37 AA.  
 XX AC ADN16159;  
 XX DT 29-JUL-2004 (first entry)  
 XX DE Human peptide hormone Amylin-analogue 25,28,29Pro-h-amylin.  
 XX

XX Human; hormone; Amylin; insulin secretion; glucagon secretion;  
 KM pancreatic; anti-inflammatory; low fat diet; pain; 25; 28;  
 KM 29Pro-h-amylin; mutant.  
 XX Homo sapiens.  
 OS Synthetic.  
 XX  
 XX Key Location/Qualifiers  
 FH Misc-difference 25 /note= "Wild-type Ala substituted by Pro"  
 FT Misc-difference 28 /note= "Wild-type Ser substituted by Pro"  
 FT Misc-difference 29 /note= "Wild-type Ser substituted by Pro"  
 FT  
 PN WO2004037168-A2.  
 PD 06-MAY-2004.  
 XX  
 XX 24-SEP-2003; 2003WO-US030224.  
 PF  
 XX 18-OCT-2002; 2002US-0419440P.  
 PR  
 XX (AMYL-) AMYLIN PHARM INC.  
 PA  
 XX Gedulin B, Young AA;  
 PI  
 XX WPI; 2004-365410/34.  
 DR  
 XX  
 XX Treating pancreaticis or relieving pain caused by pancreaticitis comprises  
 PT administering to the subject an amylin, amylin analog or an amylin  
 PT agonist.  
 PT  
 XX  
 XX Claim 9; Page; 33pp; English.  
 PS  
 XX The invention relates to treating pancreaticitis or in relieving pain  
 CC caused by pancreaticitis in a mammalian subject comprises administering to  
 CC the subject an amount of an Amylin (a peptide hormone which inhibits  
 CC insulin and glucagon secretion), Amylin analogue or an Amylin agonist,  
 CC where the Amylin agonist is not calcitonin. Also included is a method of  
 CC improving a treatment for pancreaticitis in a mammalian subject. The Amylin  
 CC analogue is 25,28,29Pro-h-amylin. The method further comprises  
 CC administering to the subject an analgesic. The regime includes a low-fat  
 CC diet. The methods are useful in treating pancreaticis or relieving pain  
 CC caused by pancreaticitis in a mammalian subject. The present sequence  
 CC represents human Amylin analogue 25,28,29Pro-h-amylin. Note: The present  
 CC sequence is not shown in the specification but was created by the indexer  
 CC using the information in claim 9 and wild-type amylin (ADN16157).  
 CC  
 XX Sequence 37 AA;  
 SQ  
 Query Match 100.0%; Score 202; DB 8; Length 37;  
 Best Local Similarity 100.0%; Pred. No. 6,4e-21;  
 Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37  
 DB 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37

RESULT 13  
 ADOS1015  
 ID ADOS1015 standard; peptide; 37 AA.  
 XX  
 XX ADO51015;  
 AC  
 XX 18-NOV-2004 (first entry)  
 DT  
 XX Human amylin agonist peptide analogue #1.  
 DE  
 XX Gastric motility; delay gastric emptying; amylin; agonist;  
 KW postprandial dumping syndrome; postprandial hyperglycemia;  
 KW

KW gastrointestinal disorder; spasm; radiological examination;  
 KM magnetic resonance imaging; diabetes; therapy; human.  
 OS Homo sapiens.  
 XX  
 XX Key Location/Qualifiers  
 FH Modified-site 37 /note= "C-terminal amide"  
 FT  
 FT  
 XX US2004097415-A1.  
 PN  
 XX 20-MAY-2004.  
 PD  
 XX 18-ANG-2003; 2003US-00643681.  
 PF  
 XX 07-SEP-1993; 93US-00118381.  
 PR 07-SEP-1994; 94US-00302069.  
 XX 22-MAY-2000; 2000US-00576062.  
 XX  
 XX (KOLT/) KOLTERMAN O G.  
 PA (YOUN/) YOUNG A A.  
 PA (RINK/) RINK T J.  
 PA (BROW/) KEATING BROWN K A.  
 XX  
 PI Kolterman OG, Young AA, Rink TJ, Keating Brown KA;  
 DR  
 XX WPI; 2004-389180/36.  
 DR  
 XX  
 XX Use of amylin agonist for reducing gastric motility, delaying gastric  
 PT emptying and for treating postprandial dumping syndrome and postprandial  
 PT hyperglycemia.  
 PT  
 XX Disclosure; SEQ ID NO 1; 35pp; English.  
 PS  
 XX The present invention is directed to novel methods for reducing gastric  
 CC motility and delaying gastric emptying which comprises the administration  
 CC of an amylin or an amylin agonist. The invention is useful for treating  
 CC postprandial dumping syndrome, postprandial hyperglycemia and reducing  
 CC gastric motility associated with gastrointestinal disorders such as spasm  
 CC or delaying gastric emptying in a mammal undergoing a gastrointestinal  
 CC diagnostic procedure such as radiological examination and magnetic  
 CC resonance imaging. The invention is also useful for lowering postprandial  
 CC blood glucose levels during treatment of diabetes. The present sequence  
 CC is human amylin agonist peptide analogue. This sequence is used in the  
 CC invention.  
 CC  
 XX Sequence 37 AA;  
 SQ  
 Query Match 100.0%; Score 202; DB 8; Length 37;  
 Best Local Similarity 100.0%; Pred. No. 6,4e-21;  
 Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37  
 DB 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37

RESULT 14  
 ADV92837  
 ID ADV92837 standard; peptide; 37 AA.  
 XX  
 XX ADV92837;  
 AC  
 XX 24-MAR-2005 (first entry)  
 DT  
 XX Amylin peptide amino acid sequence - SEQ ID 257.  
 DE  
 XX delivery mechanism; viral infections; virucide; bacterial infection;  
 KW antibacterial; amylin.  
 KW  
 XX Unidentified.  
 OS  
 XX WO2005000222-A2.  
 PN



CC pramlintide peptide, which serves as an amylin agonist peptide.  
XX  
SQ Sequence 37 AA;

Query Match 100.0%; Score 202; DB 9; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 17

AEBI7969  
ID AEBI7969 standard; peptide; 37 AA.

AC AEBI7969;

DT 08-SEP-2005 (first entry)

DE Human amylin agonist, pramlintide peptide SEQ ID NO: 47 #1.

XX Pharmaceutical; weight loss; obesity; anorectic; nutritional disorder;  
KM hyperglycemia; antidiabetic; metabolic disorder; antilipemic;  
KM diabetes mellitus; metabolic disorder; glucose regulating peptide;  
KM amylin agonist; pramlintide.

OS Homo sapiens.

XX Key Location/Qualifiers

FT Disulfide-bond 2..7

FT Modified-site 37 /note="Amidated"

XX US2005143303-A1.

XX 30-JUN-2005.

XX 18-NOV-2004; 2004US-00991597.

XX 26-DEC-2003; 2003US-0532337P.

XX (NAST-) NASTECH PHARM CO INC.

XX Quay SC, Costantino HR;

XX WPI; 2005-496434/50.

XX New transmembrane glucose-regulating peptide (GRP) formulation, useful for  
PT treating e.g. obesity, hyperglycemia, dyslipidemia and diabetes mellitus  
PT and for inducing satiety in an individual and to promote weight-loss in  
PT an individual.

PS Claim 7; SEQ ID NO 47; 55pp; English.

XX The present invention relates to pharmaceutical compositions and methods  
CC comprising at least one glucose regulating peptide (GRP) such as amylin,  
CC an amylin analog (such as pramlintide), glucagon-like peptide-1 (GLP)-1,  
CC exendin-3 or exendin-4 and one or more mucosal delivery-enhancing agents.  
CC The formulation or the method is useful for treating variety of diseases  
CC and conditions in mammalian subjects including obesity, hyperglycemia,  
CC dyslipidemia and diabetes mellitus and for inducing satiety in an  
CC individual and to promote weight-loss in an individual. The invention is  
CC also useful in protein therapy. The present sequence is the human  
CC pramlintide peptide, which serves as an amylin agonist peptide. Note: The  
CC present sequence is the SEQ ID NO: 47 which is given in the sequence  
CC listing. This sequence differs from the SEQ ID NO:47 shown on page 6 of  
CC the specification (see AEBI7970).

XX Sequence 37 AA;

Query Match 100.0%; Score 202; DB 9; Length 37;

Best Local Similarity 100.0%; Pred. No. 6.4e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 18

AAR29202  
ID AAR29202 standard; protein; 37 AA.

XX AAR29202;

AC AAR29202;

DT 25-MAR-2003 (revised)

DT 20-APR-1993 (first entry)

DE Pro(25)Val(26)Pro(28,28)-h-amylin for treating anorexia.

XX Anorexia; cachexia; adipose; amylin.

XX Homo sapiens.

XX WO9220367-A1.

XX 26-NOV-1992.

XX 23-MAY-1992; 92WO-US004357.

XX 24-MAY-1991; 91US-00704995.

XX 03-APR-1992; 92US-00862500.

XX (AMYL-) AMYLIN PHARM INC.

XX Rank TJ, Young AA;

XX WPI; 1992-415470/50.

XX Use of amylin and opt. insulin for treating anorexia - increases plasma  
PT amylin and/or insulin levels, also for treating cachexia conditions,  
PT adipose tissue deficiency etc.

XX Disclosure; Page 16; 19pp; English.

XX Treatment of a patient with anorexia or related condition comprises  
CC admin. of amylin or an analogue in an amt. sufficient to increase the  
CC amylin level in the plasma of the patient. The pref. amylin analogues are  
CC given in AAR29197-222. Treating a patient deficient in adipose tissue

CC comprises admin. of amylin or an analogue and/or insulin in an amt. and  
CC ratio sufficient to increase adipose tissue. Typical dosage units contain  
CC 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-  
CC 2003 to correct FN field.)

XX Sequence 37 AA;

XX Query Match 99.5%; Score 201; DB 2; Length 37;

XX Best Local Similarity 97.3%; Pred. No. 8.9e-21;

XX Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 19

AAR29212  
ID AAR29212 standard; protein; 37 AA.

XX AAR29212;

AC AAR29212;

DT 25-MAR-2003 (revised)

DT 20-APR-1993 (first entry)

DE Ile(17)Pro(25,28,29)-h-amylin for treating anorexia.  
 XX Anorexia; cachexia; adipose; amylin.  
 XX Homo sapiens.  
 OS MO9220367-A1.  
 PN 26-NOV-1992.  
 PD  
 XX  
 PF 23-MAY-1992; 92WO-US004357.  
 XX  
 PR 24-MAY-1991; 91US-00704995.  
 PR 03-APR-1992; 92US-00862500.  
 XX  
 PA (AMYL-) AMYLIN PHARM INC.  
 XX  
 PI Rink TJ, Young AA;  
 DR WPI; 1992-415470/50.  
 XX  
 PT Use of amylin and opt. insulin for treating anorexia - increases plasma  
 PT amylin and/or insulin levels, also for treating cachexia conditions,  
 PT adipose tissue deficiency etc.  
 XX  
 XX Disclosure; Page 16; 19pp; English.  
 PS  
 CC Treatment of a patient with anorexia or related condition comprises  
 CC admin. of amylin or an analogue in an amt. sufficient to increase the  
 CC amylin level in the plasma of the patient. The pref. amylin analogues are  
 CC given in AAR29197-222. Treating a patient deficient in adipose tissue  
 CC comprises admin. of amylin or an analogue and/or insulin in an amt. and  
 CC ratio sufficient to increase adipose tissue. Typical dosage units contain  
 CC 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-  
 CC 2003 to correct PN field.)  
 XX  
 SQ Sequence 37 AA;  
 QY  
 Query Match 99.5%; Score 201; DB 2; Length 37;  
 Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
 Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 Db 1 KCNTATCATORLANFLVHSSNNFGPILPPTVGSNTY 37  
 1 KCNTATCATORLANFLVHSSNNFGPILPPTVGSNTY 37  
 RESULT 20  
 AAR37780  
 ID AAR37780 standard; peptide; 37 AA.  
 XX  
 AC AAR37780;  
 XX  
 XX 25-MAR-2003 (revised)  
 DT 07-SEP-1993 (first entry)  
 XX  
 DE Pro25, Val26, Pro28, Pro29 human amylin analogue.  
 XX  
 KW Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;  
 KW hyperglycaemic agent.  
 OS Homo sapiens.  
 XX  
 XX Key Location/Qualifiers  
 FH Disulfide-bond 2..7  
 FT Misc-difference 25  
 FT Misc-difference 26 /note= "Pro replaces wild-type Ala"  
 FT Misc-difference 28 /note= "Val replaces wild-type Ile"  
 FT Misc-difference 29 /note= "Pro replaces wild-type Ser"  
 FT Misc-difference 29 /note= "Pro replaces wild-type Ser"

FT Modified-site 37 /note= "amidated"  
 FT  
 XX  
 PN WO9310146-A1.  
 XX  
 PD 27-MAY-1993.  
 XX  
 PF 19-NOV-1992; 92WO-US009842.  
 XX  
 PR 19-NOV-1991; 91US-00794266.  
 XX  
 PA (AMYL-) AMYLIN PHARM INC.  
 XX  
 PI Gaeta LSL, Jones H, Albrecht E;  
 DR WPI; 1993-182488/22.  
 XX  
 PT New amylin agonist peptide(s) - used for treatment and prevention of  
 PT hypoglycaemia and diabetes mellitus.  
 XX  
 PS Example 2; Fig 1 and Page 16; 43pp; English.  
 CC  
 CC This peptide is an example of amylin agonists of the invention which can  
 CC be used as hyperglycaemics. The peptide is an analogue of human amylin  
 CC which mimics the effects of the wild-type hormone. Preferred peptides are  
 CC used in admixture with insulin for the treatment of diabetes mellitus or  
 CC with glucagon for the treatment of hypoglycaemic conditions. See AAR37779  
 CC field.)  
 CC  
 CC R37795 and AAR38809-R38826. (Updated on 25-MAR-2003 to correct PN  
 CC field.)  
 XX  
 SQ Sequence 37 AA;  
 QY  
 Query Match 99.5%; Score 201; DB 2; Length 37;  
 Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
 Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 Db 1 KCNTATCATORLANFLVHSSNNFGPILPPTVGSNTY 37  
 1 KCNTATCATORLANFLVHSSNNFGPILPPTVGSNTY 37  
 RESULT 21  
 AAR38816  
 ID AAR38816 standard; peptide; 37 AA.  
 XX  
 AC AAR38816;  
 XX  
 XX 25-MAR-2003 (revised)  
 DT 07-SEP-1993 (first entry)  
 XX  
 DE Ile17, Pro25, Pro28, Pro29-human amylin analogue.  
 XX  
 KW Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;  
 KW hyperglycaemic agent.  
 OS Homo sapiens.  
 XX  
 XX Key Location/Qualifiers  
 FH Disulfide-bond 2..7  
 FT Misc-difference 17 /note= "Ile replaces wild-type Val"  
 FT Misc-difference 25 /note= "Pro replaces wild-type Ala"  
 FT Misc-difference 28 /note= "Pro replaces wild-type Ser"  
 FT Misc-difference 29 /note= "Pro replaces wild-type Ser"  
 FT Modified-site 37 /note= "amidated"  
 FT  
 XX  
 PN WO9310146-A1.  
 XX  
 PD 27-MAY-1993.

Qy	Query Match	99.5%	Score 201;	DB 2;	Length 37;
db	Best Local Similarity	97.3%;	Pred. No. 8.9e-21;		
	Matches 36;	Conservative 1;	Mismatches 0;	Indels 0;	Gaps 0;
1	KCNTATCATCATORLANFLVHSSNNFGCGLPPTVNGSNTY	37			
1	KCNTATCATCATORLANFLVHSSNNFGCGLPPTVNGSNTY	37			

XX	WP1; 1999-458254/38.
DR	
XX	Stabilized liquid formulation for treatment of insulin-dependent diabetes
PT	mellitus.
XX	
PS	Disclosure; Page; 71pp; English.
XX	
CC	This sequence represents a human amylin analogue, that acts as a amylin
CC	agonist. The invention relates to a liquid pharmaceutical formulation
CC	(A), that contains (wt./vol.): 0.01-0.5% amylin agonist (I); 1-10%
CC	carbohydrate or polyol (II); and 0.02-0.5% acetate, phosphate, citrate or
CC	glutamate buffer (III); and has pH 3-6. (A) are used, in conjunction with
CC	insulin, for treatment of diabetes, specifically to reduce post-prandial
CC	increases in glucose levels of the blood. In these formulations, (I) is
CC	stabilised, especially against deamidation and peptide bond hydrolysis
CC	for up to 4 years at 5 degrees C and 30 days at 30 degrees C, without
CC	addition of a separate stabiliser. They also retain short-term (up to 24
CC	hr) stability when combined with insulin, allowing both agents to be
CC	administered together, reducing the number of injections required. Note:
CC	This sequence was created by the indexer from information given in the
CC	specification
XX	
SQ	Sequence 37 AA;
Query Match	99.5%; Score 201; DB 2; Length 37;
Best Local Similarity	97.3%; Pred. No. 8.9e-21;
Matches	36; Conservative 1; Mismatches 0; Indels 0; Gaps 0
QY	1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37
DB	1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37
RESULT 23	
ID	AAV22449
AAV22449	standard; peptide; 37 AA.
XX	
AC	AAV22449;
XX	
DT	28-SEP-1999 (first entry)
XX	
DE	11e17, Pro25, Pro28, Pro29 amylin analogue.
XX	
KW	Amylin agonist; human; insulin; diabetes; post-prandial glucose level;
XX	therapy; mutcin.
XX	
OS	Homo sapiens.
OS	Synthetic.
XX	
FH	Key
FT	Disulfide-bond 2..7
FT	Misc-difference 17
FT	/label= V171
FT	Misc-difference 25
FT	/label= A25P
FT	Misc-difference 28
FT	/label= S28P
FT	Misc-difference 29
FT	/label= S29P
XX	
XX	
XX	WO9934822-A1.
XX	
XX	15-JUL-1999.
XX	
XX	09-JAN-1998; 98WO-US000288.
XX	
XX	09-JAN-1998; 98WO-US000288.
XX	
XX	(AMYL-) AMYLIN PHARM INC.
XX	
XX	L'italian J, Musunuri S, Ruby C;
XX	

DR WPI, 1999-458254/38.  
XX Stabilised liquid formulation for treatment of insulin-dependent diabetes  
PT mellitus.  
XX  
XX Disclosure, Page, 71pp; English.  
XX  
XX This sequence represents a human amylin analogue, that acts as a amylin  
CC agonist. The invention relates to a liquid pharmaceutical formulation  
CC (A), that contains (wt./vol.): 0.01-0.5% amylin agonist (I); 1-10%  
CC carbohydrate or polyol (II); and 0.02-0.5% acetate, phosphate, citrate or  
CC glutamate buffer (III); and has pH 3-6. (A) are used, in conjunction with  
CC insulin, for treatment of diabetes, specifically to reduce post-prandial  
CC increases in glucose levels of the blood. In these formulations, (I) is  
CC stabilised, especially against deamidation and peptide bond hydrolysis  
CC for up to 4 years at 5 degrees C and 30 days at 30 degrees C, without  
CC addition of a separate stabiliser. They also retain short-term (up to 24  
CC hr) stability when combined with insulin, allowing both agents to be  
CC administered together, reducing the number of injections required. Note:  
CC This sequence was created by the indexer from information given in the  
CC specification  
XX  
SQ Sequence 37 AA:  
  
Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
  
RESULT 24  
ID AAW90146 standard; peptide; 37 AA.  
AC AAW90146;  
XX  
XX 15-MAR-1999 (first entry)  
XX  
XX Human amylin agonist peptide 25-Pro26-Val28,29-Pro-amylin.  
DE  
XX Amylin; human; agonist; gastritis; gastric ulceration; treatment;  
XX non-steroidal anti-inflammatory agent; NSAID; intravenous; subcutaneous;  
XX pain; fever; inflammation; arthritis; hypercoagulation.  
XX  
XX Homo sapiens.  
OS Synthetic.  
XX  
XX Key Location/Qualifiers  
XX FT Disulfide-bond 2..7  
XX  
XX WO9850059-A1.  
XX  
XX 12-NOV-1998.  
XX  
XX 06-MAY-1998; 98WO-US009089.  
XX  
XX 06-MAY-1997; 97US-00851965.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX Young A, Gedulin B, Beynon GW;  
XX  
XX WPI, 1999-059652/05.  
XX  
XX Method for treating or preventing gastritis - comprises administering  
XX amylin or amylin agonist, except calcitonin.  
XX  
XX Claim 6; Page 42; 48pp; English.  
XX  
XX This invention relates to a method for treating or preventing gastritis

CC or gastric ulceration which comprises administering amylin or an amylin  
CC agonist. Amylin administration is not carried out intra-  
CC cerebroventricularly. The specification describes a method for treating  
CC or preventing a condition for which a non-steroidal anti-inflammatory  
CC agent (NSAID) is indicated, comprising administering amylin or amylin  
CC agonist, which is not calcitonin, together with NSAID and also a  
CC composition comprising an amylin or an amylin agonist or their salts,  
CC except calcitonin and a NSAID in a carrier. The amylin composition is  
CC used to treat humans by administering it subcutaneously, intravenously or  
CC by nasal, oral, pulmonary, transdermal and buccal routes. The method is  
CC also used to treat pain, fever, inflammation, arthritis, hypercoagulation  
CC and other condition where an NSAID would be indicated. The present  
CC sequence is an example of an agonist used in the method  
XX  
SQ Sequence 37 AA:  
  
Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
  
RESULT 25  
ID AAW90154 standard; peptide; 37 AA.  
AC AAW90154;  
XX  
XX 15-MAR-1999 (first entry)  
XX  
XX Human amylin agonist peptide 17-Ile25,28,29-Pro-amylin.  
DE  
XX  
XX Amylin; human; agonist; gastritis; gastric ulceration; treatment;  
XX non-steroidal anti-inflammatory agent; NSAID; intravenous; subcutaneous;  
XX pain; fever; inflammation; arthritis; hypercoagulation.  
XX  
XX Homo sapiens.  
OS Synthetic.  
XX  
XX Key Location/Qualifiers  
XX FT Disulfide-bond 2..7  
XX  
XX WO9850059-A1.  
XX  
XX 12-NOV-1998.  
XX  
XX 06-MAY-1998; 98WO-US009089.  
XX  
XX 06-MAY-1997; 97US-00851965.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX Young A, Gedulin B, Beynon GW;  
XX  
XX WPI, 1999-059652/05.  
XX  
XX Method for treating or preventing gastritis - comprises administering  
XX amylin or amylin agonist, except calcitonin.  
XX  
XX Claim 6; Page 42; 48pp; English.  
XX  
XX This invention relates to a method for treating or preventing gastritis  
CC or gastric ulceration which comprises administering amylin or an amylin  
CC agonist. Amylin administration is not carried out intra-  
CC cerebroventricularly. The specification describes a method for treating  
CC or preventing a condition for which a non-steroidal anti-inflammatory  
CC agent (NSAID) is indicated, comprising administering amylin or amylin  
CC agonist, which is not calcitonin, together with NSAID and also a  
CC composition comprising an amylin or an amylin agonist or their salts,  
CC except calcitonin and a NSAID in a carrier. The amylin composition is



CC used to treat humans by administering it subcutaneously, intravenously or  
CC by nasal, oral, pulmonary, transdermal and buccal routes. The method is  
CC also used to treat pain, fever, inflammation, arthritis, hypercoagulation  
CC and other condition where an NSAID would be indicated. The present  
CC sequence is an example of an agonist used in the method

XX Sequence 37 AA;

Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8,9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37

RESULT 26  
AAB18578  
XX AAB18578 standard; peptide; 37 AA.  
XX  
XX AAB18578;  
XX  
XX 15-JAN-2001 (first entry)  
XX  
XX

DE Amino acid sequence of an amylin agonist analogue compound.

XX  
XX Amylin agonist; amylin; gastric motility; gastric emptying;  
XX postprandial dumping syndrome; postprandial hyperglycemia;  
XX gastrointestinal diagnostic procedure; gastrointestinal disorder; spasm;  
XX acute diverticulitis; biliary tract disorder; sphincter of oddi disorder.

OS Synthetic.

XX  
XX Key Location/Qualifiers  
FH Disulfide-bond 2..7  
FT Modified-site 37  
FT /note= "amidated residue"

XX US6114304-A.

XX 05-SEP-2000.

XX 07-SEP-1994; 94US-00302069.

XX 07-SEP-1993; 93US-00118381.

XX (AMYL-) AMYLIN PHARM INC.

XX Young AA, Rink TJ, Brown KAK, Kolterman OG;

XX WPI, 2000-601336/57.

XX Treating gastrointestinal disorder e.g. spasm by reducing gastric  
XX motility or delaying gastric emptying, postprandial dumping syndrome or  
XX postprandial hyperglycemia, by administering amylin or amylin agonist.

XX Disclosure; Col 39-40; 50pp; English.

XX The present sequence represents an amylin agonist analogue compound.  
XX Amylin or amylin agonists are administered for reducing gastric motility  
XX or delaying gastric emptying, and for treating postprandial dumping  
XX syndrome or postprandial hyperglycemia, by inducing amylin activity, in a  
XX mammal. The peptides are used to reduce gastric motility or for delaying  
XX gastric emptying in a mammal undergoing gastrointestinal diagnostic  
XX procedures, such as radiological examination or magnetic resonance  
XX imaging. They are also used for reducing gastric motility in  
XX gastrointestinal disorder, especially spasm, which is associated with a  
XX sphincter of oddi. They are also used to treat postprandial dumping  
XX syndrome or postprandial hyperglycemia

XX Sequence 37 AA;

Query Match 99.5%; Score 201; DB 3; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8,9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37

RESULT 27  
AAB18589  
XX AAB18589 standard; peptide; 37 AA.  
XX  
XX AAB18589;  
XX  
XX 15-JAN-2001 (first entry)  
XX  
XX

DE Amino acid sequence of an amylin agonist analogue compound.

XX  
XX Amylin agonist; amylin; gastric motility; gastric emptying;  
XX postprandial dumping syndrome; postprandial hyperglycemia;  
XX gastrointestinal diagnostic procedure; gastrointestinal disorder; spasm;  
XX acute diverticulitis; biliary tract disorder; sphincter of oddi disorder.

OS Synthetic.

XX  
XX Key Location/Qualifiers  
FH Disulfide-bond 2..7  
FT Modified-site 37  
FT /note= "amidated residue"

XX US6114304-A.

XX 05-SEP-2000.

XX 07-SEP-1994; 94US-00302069.

XX 07-SEP-1993; 93US-00118381.

XX (AMYL-) AMYLIN PHARM INC.

XX Young AA, Rink TJ, Brown KAK, Kolterman OG;

XX WPI, 2000-601336/57.

XX Treating gastrointestinal disorder e.g. spasm by reducing gastric  
XX motility or delaying gastric emptying, postprandial dumping syndrome or  
XX postprandial hyperglycemia, by administering amylin or amylin agonist.

XX Disclosure; Col 47-48; 50pp; English.

XX The present sequence represents an amylin agonist analogue compound.  
XX Amylin or amylin agonists are administered for reducing gastric motility  
XX or delaying gastric emptying, and for treating postprandial dumping  
XX syndrome or postprandial hyperglycemia, by inducing amylin activity, in a  
XX mammal. The peptides are used to reduce gastric motility or for delaying  
XX gastric emptying in a mammal undergoing gastrointestinal diagnostic  
XX procedures, such as radiological examination or magnetic resonance  
XX imaging. They are also used for reducing gastric motility in  
XX gastrointestinal disorder, especially spasm, which is associated with a  
XX disorder of acute diverticulitis or disorders of biliary tract or  
XX sphincter of oddi. They are also used to treat postprandial dumping  
XX syndrome or postprandial hyperglycemia

XX Sequence 37 AA;

Query Match 99.5%; Score 201; DB 3; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8,9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37  
1 KCNTATCATQRLANFLVHSSNNGFPIPTNVGSNTY 37

DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37

RESULT 28  
ABB05492  
ID ABB05492 standard; peptide; 37 AA.  
XX  
XX ABB05492;  
XX  
DT 19-APR-2002 (first entry)  
XX  
DE Human amylin agonist 25ProGVal28, 29Pro-h-amylin.  
XX  
XX Human; amylin agonist; amylin peptide analogue; acetate; carbohydrate;  
KW liquid pharmaceutical formulation; polyhydric alcohol; phosphate;  
KW citrate; glutamate; buffer; antidiabetic; type II diabetes.  
XX  
XX Homo sapiens.  
OS Synthetic.  
OS  
PN US2001043934-A1.  
XX  
PD 22-NOV-2001.  
XX  
PF 09-JAN-1998; 98US-00005262.  
XX  
PR 08-JAN-1997; 97US-0035140P.  
XX  
XX (LIT/)/ L'ITALIEN J.  
PA (MUSU/) MUSUNURI S.  
PA (RUBY/) RUBY K.  
XX  
XX L'Italien J, Musunuri S, Ruby K;  
PI  
XX  
XX WPI; 2002-163554/21.  
DR  
PT New pharmaceutical formulation useful for treating patients with type II  
PT diabetes comprises amylin agonist, carbohydrate or polyhydric alcohol and  
PT a buffer.  
XX  
XX  
XX Disclosure; Page; 19pp; English.  
XX  
XX The present invention describes a liquid pharmaceutical formulation (A)  
CC comprising (w/vol%) an amylin agonist (0.01 - 0.5), a carbohydrate or a  
CC polyhydric alcohol (1 - 10) and an acetate, phosphate, citrate or  
CC glutamate buffer (0.02 - 0.5) having a pH of 3 - 6. Also described is a  
CC commercial package containing the liquid pharmaceutical formulation (A).  
CC The package comprises a borosilicate glass vial having an open end, a  
CC stopper for multise compatible with the amylin and/or amylin agonist  
CC fixed in the open end of the vial. The package also comprises a  
CC cartridge for use in a pen injector. (A) has antidiabetic activity and  
CC can be used in the treatment of patients with type II diabetes. The  
CC formulation comprises amylin agonist which is biologically active, has a  
CC reduced tendency to form aggregates in water or at a pressure of greater  
CC than 2 psi and has a reduced tendency to precipitate in the presence of  
CC NaCl compared to human amylin. The formulation maintains stability upon  
CC storage under refrigerated or room-temperature conditions. The  
CC formulation retains short-term mixing compatibility with insulin and  
CC results in improved stability of the hormone and the patients no longer  
CC need to refrigerate the vial of insulin in use. The present sequence  
CC represents a human amylin peptide analogue, which can be used as an  
CC amylin agonist in the present invention. N.B. The present sequence is not  
CC given in the present specification but is derived from the 37 amino acid  
CC human amylin as stated in the invention  
XX  
XX  
SQ Sequence 37 AA;

Query Match 99.5%; Score 201; DB 5; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37

DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37

RESULT 29  
ABB05504  
ID ABB05504 standard; peptide; 37 AA.  
XX  
XX ABB05504;  
XX  
DT 19-APR-2002 (first entry)  
XX  
DE Human amylin agonist 17Ile25, 28, 29Pro-h-amylin.  
XX  
XX Human; amylin agonist; amylin peptide analogue; acetate; carbohydrate;  
KW liquid pharmaceutical formulation; polyhydric alcohol; phosphate;  
KW citrate; glutamate; buffer; antidiabetic; type II diabetes.  
XX  
XX Homo sapiens.  
OS Synthetic.  
OS  
PN US2001043934-A1.  
XX  
PD 22-NOV-2001.  
XX  
PF 09-JAN-1998; 98US-00005262.  
XX  
PR 08-JAN-1997; 97US-0035140P.  
XX  
XX (LIT/)/ L'ITALIEN J.  
PA (MUSU/) MUSUNURI S.  
PA (RUBY/) RUBY K.  
XX  
XX L'Italien J, Musunuri S, Ruby K;  
PI  
XX  
XX WPI; 2002-163554/21.  
DR  
PT New pharmaceutical formulation useful for treating patients with type II  
PT diabetes comprises amylin agonist, carbohydrate or polyhydric alcohol and  
PT a buffer.  
XX  
XX  
XX Disclosure; Page; 19pp; English.  
XX  
XX The present invention describes a liquid pharmaceutical formulation (A)  
CC comprising (w/vol%) an amylin agonist (0.01 - 0.5), a carbohydrate or a  
CC polyhydric alcohol (1 - 10) and an acetate, phosphate, citrate or  
CC glutamate buffer (0.02 - 0.5) having a pH of 3 - 6. Also described is a  
CC commercial package containing the liquid pharmaceutical formulation (A).  
CC The package comprises a borosilicate glass vial having an open end, a  
CC stopper for multise compatible with the amylin and/or amylin agonist  
CC fixed in the open end of the vial and an aluminum band to retain the  
CC stopper in the far end of the vial. The package also comprises a  
CC cartridge for use in a pen injector. (A) has antidiabetic activity and  
CC can be used in the treatment of patients with type II diabetes. The  
CC formulation comprises amylin agonist which is biologically active, has a  
CC reduced tendency to form aggregates in water or at a pressure of greater  
CC than 2 psi and has a reduced tendency to precipitate in the presence of  
CC NaCl compared to human amylin. The formulation maintains stability upon  
CC storage under refrigerated or room-temperature conditions. The  
CC formulation retains short-term mixing compatibility with insulin and  
CC results in improved stability of the hormone and the patients no longer  
CC need to refrigerate the vial of insulin in use. The present sequence  
CC represents a human amylin peptide analogue, which can be used as an  
CC amylin agonist in the present invention. N.B. The present sequence is not  
CC given in the present specification but is derived from the 37 amino acid  
CC human amylin as stated in the invention  
XX  
XX  
SQ Sequence 37 AA;

Query Match 99.5%; Score 201; DB 5; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.9e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;





DR WPI; 2005-075434/08.

PT Pharmaceutical composition for transnasal administration of a bioactive peptide/protein of interest, e.g. glucagon-like peptide-1, comprises the peptide/protein of interest, a cationic polyamino acid, and a compatible buffer.

PS Disclosure; SEQ ID NO 260; 64pp; English.

The invention comprises a pharmaceutical composition for transmucosal administration of a bioactive peptide/protein (e.g. exendin, PYY, GLP-1 or amylin peptide/protein) of interest. The composition of the invention is useful for the transmucosal administration of a bioactive peptide or protein and is useful for treating or preventing viral or bacterial diseases in humans. The present amino acid sequence represents an amylin peptide that is used in the exemplification of the invention.

**SQ Sequence 37 AA;**

```
Query Match      99.5%;   Score 201;   DB 9;   Length 37;
Best Local Simlarity 97.3%;   Pred. No. 8.9e-21;
Matches 36;   Conservative 1;   Mismatches 0;   Indels 0;   Gaps 0;
```

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
|||||:|||||  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

```
RESULT 35
AEBI7939
ID AEBI7939 standard; peptide; 37 AA.
```

AC AEB17939;

DT 08-SEP-2005 (first entry)

Human amylin agonist, pramlintide peptide SEQ ID NO: 17.

KM Pharmacological: weight loss; obesity; anorectic; nutritional disorder  
KM hyperglycemia; antidiabetic; metabolic disorder; antilipemic;  
KM diabetes mellitus; metabolic disorder; glucose regulating peptide;  
KM amylin agonist; pramlintide.

OS Homo sapiens.

	Location/Qualifiers
FH	Key
FT	Disulfide-bond
FT	Modified-site
FT	37
	/note= "Amidated"

PN US2005143303-A1.

PD 30-JUN-2005.

PF 18-NOV-2004; 2004US-00991597.

PR 26-DEC-2003; 2003US-0532337P.

PA (NAST-) NASTECH PHARM CO INC.

Quay SC, Costantino HR;

DR WPI; 2005-496434/50.

PT New transmembrane glucose-regulating peptide (GRP) formulation, useful for  
PT treating e.g. obesity, hyperglycemia, dyslipidemia and diabetes mellitus  
PT and for inducing satiety in an individual and to promote weight-loss in  
PT an individual.

PS Claim 7; SEQ ID NO 17; 55pp; English.

CC The present invention relates to pharmaceutical compositions and methods  
CC comprising at least one glucose regulating peptide (GRP) such as amylin,

CC an amylin analog (such as pramlintide), glucagon-like peptide-1 (GLP-1),  
CC extendin-3 or extendin-4 and one or more mucosal delivery-enhancing agents  
CC The formulation or the method is useful for treating variety of diseases  
CC and conditions in mammalian subjects including obesity, hyperglycemia,  
CC dyslipidemia and diabetes mellitus and for inducing satiety in an  
CC individual and to promote weight-loss in an individual. The invention is  
CC also useful in protein therapy. The present sequence is the human  
CC pramlintide peptide, which serves as an amylin agonist peptide.

**SQ** Sequence 37 AA;

Query Match	99.5%	Score 201	DB 9	Length 37
Best Local Similarity	97.3%	Pred. No. 8	9e-21	
Matches 36	Conservative 1	Mismatches 0	Indels 0	Gaps 0

QY	1	KCNTATCATQRLANFLVHSSNNGFPIILPTTVG	SNTY	37
			:	
Db	1	KCNTATCATQRLANFLVHSSNNGFVLPPTTVG	SNTY	37

RESULT 36	
AAW22541	
ID AAW22541	standard; peptide; 37 AA

AC AAW22541;

DT 12-OCT-1997 (first entry)

DE Amylin agonist peptide 11-Lys-25,28,29-Pro-h-amylin

KW appetite regulation; amylin agonist; hybrid; CCK; cholecystokinin

OS Synthetic.

	Location/Qualifiers
FH Key	2. .7
FT Disulfide-bond	37
FT Modified-site	/notes= "Tyr-NH2"
FT	

PN W09640196-A1.

PD 19-DEC-1996

PF 06-JUN-1996; 96WO-US009937.

PR 07-JUN-1995; 95US-00477727.

PA (AMYL-) AMYLIN PHARM INC.

PI Rink TJ, Young AA, Beeley NR, Prickett KS;

DR WPI; 1997-051885/05.

PT Compositions comprising amylin and cholecystokinin agonists - useful for  
PT reducing food intake, suppressing appetite and controlling body weight.  
XX

XS Claim 20,49; Page 46,54; 66pp; English.

The patent discloses a composition which can be used to reduce or suppress food intake, control appetite or control body weight in a mammal, comprising an amylin agonist and a cholecystokinin (CCK) agonist admixed in a form suitable for therapeutic administration. Also disclosed are new hybrid peptides comprising an amylin agonist peptide and a CCK agonist peptide covalently linked e.g. by the group -R1-R2-R3-R4-R5- where R1 = CONH(CH2)n, COO(CH2)n or CO(CH2)n; R2 = OCO(CH2)n, NHCO(CH2)n, OCOCH4 (ortho, meta or para linked), COOCH4 or NHCOCH4 (both ortho, meta or para linked/substituted), CONHCH4NH (ortho, meta or para substituted), O-X or NH-X; R3 = CH2, CCF4, CO, CS or CNH; R4 = O or NH; R5 = (CH2)nmHCO, (CH2)nOCO or (CH2)nCO; n = 1-6; and X = any amino acid linked via its carboxyl group. Administration of amylin and CCK agonists in conjunction produces a greater effect than either administered alone; e.g. 0.1 microgram/kg of each peptide causes a substantial reduction of food intake about equivalent to that seen with 100 microgram/kg of either

CC peptide alone. The present sequence represents a preferred amylin agonist  
 CC peptide which can be used as a component of the hybrid peptide  
 XX  
 SQ Sequence 37 AA;

Query Match 98.5%; Score 199; DB 2; Length 37;  
 Best Local Similarity 97.3%; Pred. No. 1.7e-20;  
 Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
 |||||  
 DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 37  
 AAR29204  
 ID AAR29204 standard; protein; 36 AA.  
 XX  
 AC AAR29204;

XX 25-MAR-2003 (revised)  
 DT 20-APR-1993 (first entry)  
 XX

DE Des-Lys(1)Pro(25,28,29)-h-amylin for treating anorexia.

KM Anorexia; cachexia; adipose; amylin.

OS Homo sapiens.

XX MO9220367-A1.

XX 26-NOV-1992.

XX 23-MAY-1992; 92WO-US004357.

XX 24-MAY-1991; 91US-00704995.

PR 03-APR-1992; 92US-00862500.

PA (AMYL-) AMYLIN PHARM INC.

PI Rink Tü, Young AA;

XX WPI; 1992-415470/50.

XX Use of amylin and opt. insulin for treating anorexia - increases plasma  
 PT amylin and/or insulin levels, also for treating cachexia conditions,  
 PT adipose tissue deficiency etc.  
 XX

PS Disclosure; Page 16; 19pp; English.

XX Treatment of a patient with anorexia or related condition comprises  
 CC admin. of amylin or an analogue in an amt. sufficient to increase the  
 CC amylin level in the plasma of the patient. The pref. amylin analogues are  
 CC given in AAR29197-222. Treating a patient deficient in adipose tissue  
 CC comprises admin. of amylin or an analogue and/or insulin in an amt. and  
 CC ratio sufficient to increase adipose tissue. Typical dosage units contain  
 CC 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-  
 CC 2003 to correct PN field.)  
 XX

SQ Sequence 36 AA;

Query Match 97.5%; Score 197; DB 2; Length 36;  
 Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
 |||||  
 DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 38  
 AAR37793  
 ID AAR37793 standard; peptide; 36 AA.

XX AAR37793;

AC 25-MAR-2003 (revised)

XX 07-SEP-1993 (first entry)

DE (Des-Lys1), Pro25, Pro28, Pro29 human amylin analogue.

KM Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;  
 KW hyperglycaemic agent.

XX Homo sapiens.

OS Key Location/Qualifiers

XX Disulfide-bond 1..6

FT Misc-difference 24 /note= "Pro replaces wild-type Ala"

FT Misc-difference 27 /note= "Pro replaces wild-type Ser"

FT Misc-difference 28 /note= "Pro replaces wild-type Ser"

FT Modified-site 36 /note= "amidated"

XX WO9310146-A1.

XX 27-MAY-1993.

XX 19-NOV-1992; 92WO-US009842.

XX 19-NOV-1991; 91US-00794266.

XX (AMYL-) AMYLIN PHARM INC.

PI Gaeta LSL, Jones H, Albrecht E;

XX WPI; 1993-182488/22.

XX New amylin agonist peptide(s) - used for treatment and prevention of  
 PT hypoglycaemia and diabetes mellitus.  
 XX

PS Claim 44; Fig 1 and Page 23; 43pp; English.

XX This peptide is an example of amylin agonists of the invention which can  
 CC be used as hyperglycaemics. The peptide is an analogue of human amylin  
 CC which mimics the effects of the wild-type hormone. Preferred peptides are  
 CC used in admixture with insulin for the treatment of diabetes mellitus or  
 CC with glucagon for the treatment of hypoglycaemic conditions. See AAR3779  
 CC -R37795 and AAR38809-R38826. (Updated on 25-MAR-2003 to correct PN  
 CC field.)  
 XX

SQ Sequence 36 AA;

Query Match 97.5%; Score 197; DB 2; Length 36;  
 Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
 |||||  
 DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 39  
 AAY22442  
 ID AAY22442 standard; peptide; 36 AA.

AC AAY22442;

DT 28-SEP-1999 (first entry)

DE Des-Lys1, Pro25, Pro28, Pro29 amylin analogue.

KW Amylin agonist; human; insulin; diabetes; post-prandial glucose level;

KW therapy; mutein.  
XX  
OS Homo sapiens.  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Disulfide-bond 1..6  
FT Misc-difference 24  
FT Misc-difference 27 /label= A25P  
FT Misc-difference 27 /label= S28P  
FT Misc-difference 28 /label= S29P  
XX  
XX WO9934822-A1.  
XX  
XX 15-JUL-1999.  
XX  
XX 09-JAN-1998; 98WO-US000288.  
XX  
XX 09-JAN-1998; 98WO-US000288.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX L'italian J, Musunuri S, Ruby C;  
XX WPI; 1999-458254/38.  
XX  
XX Stabilized liquid formulation for treatment of insulin-dependent diabetes mellitus.  
XX  
XX Disclosure; Page; 71pp; English.  
XX  
XX This sequence represents a human amylin analogue, that acts as a amylin agonist. The invention relates to a liquid pharmaceutical formulation (A), that contains (wt./vol.): 0.01-0.5% amylin agonist (I); 1-10% carbohydrate or polyol (II); and 0.02-0.5% acetate, phosphate, citrate or glutamate buffer (III); and has pH 3-6. (A) are used, in conjunction with insulin, for treatment of diabetes, specifically to reduce post-prandial increases in glucose levels of the blood. In these formulations, (I) is stabilized, especially against deamidation and peptide bond hydrolysis for up to 4 years at 5 degrees C and 30 days at 30 degrees C, without addition of a separate stabiliser. They also retain short-term (up to 24 hr) stability when combined with insulin, allowing both agents to be administered together, reducing the number of injections required. Note: This sequence was created by the indexer from information given in the specification  
XX  
XX Sequence 36 AA;  
SQ  
Query Match 97.5%; Score 197; DB 2; Length 36;  
Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36  
RESULT 40  
AAW90145  
ID AAW90145 standard; peptide; 36 AA.  
XX  
XX AAW90145;  
XX  
XX 15-MAR-1999 (first entry)  
XX  
XX Human amylin agonist peptide des-1-Lys25,28,29-Pro-amylin.  
XX  
XX Amylin; human; agonist; gastritis; gastric ulceration; treatment;  
KW non-steroidal anti-inflammatory agent; NSAID; intravenous; subcutaneous;  
KW pain; fever; inflammation; arthritis; hypercoagulation.  
XX

OS Homo sapiens.  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Disulfide-bond 1..6  
XX  
XX WO9850059-A1.  
XX  
XX 12-NOV-1998.  
XX  
XX 06-MAY-1998; 98WO-US0009089.  
XX  
XX 06-MAY-1997; 97US-00851965.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX Young A, Gedulin B, Beynon GW,  
XX WPI; 1999-059652/05.  
XX  
XX Method for treating or preventing gastritis - comprises administering amylin or amylin agonist, except calcitonin.  
XX  
XX Claim 6; Page 42; 48pp; English.  
XX  
XX This invention relates to a method for treating or preventing gastritis or gastric ulceration which comprises administering amylin or an amylin agonist. Amylin administration is not carried out intra-cerebroventricularly. The specification describes a method for treating or preventing a condition for which a non-steroidal anti-inflammatory agent (NSAID) is indicated, comprising administering amylin or amylin agonist, which is not calcitonin, together with NSAID and also a composition comprising an amylin or an amylin agonist or their salts, except calcitonin and a NSAID in a carrier. The amylin composition is used to treat humans by administering it subcutaneously, intravenously or by nasal, oral, pulmonary, transdermal and buccal routes. The method is also used to treat pain, fever, inflammation, arthritis, hypercoagulation and other condition where an NSAID would be indicated. The present sequence is an example of an agonist used in the method  
XX  
XX Sequence 36 AA;  
SQ  
Query Match 97.5%; Score 197; DB 2; Length 36;  
Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36  
RESULT 41  
AAB18581  
ID AAB18581 standard; peptide; 36 AA.  
XX  
XX AAB18581;  
XX  
XX 15-JAN-2001 (first entry)  
XX  
XX Amino acid sequence of an amylin agonist analogue compound.  
XX  
XX Amylin agonist; amylin; gastric motility; gastric emptying;  
KW postprandial dumping syndrome; postprandial hyperglycemia;  
KW gastrointestinal diagnostic procedure; gastrointestinal disorder; spasm;  
KW acute diverticulitis; biliary tract disorder; sphincter of oddi disorder.  
XX  
XX Synthetic.  
XX  
XX Key Location/Qualifiers  
FH Disulfide-bond 1..6  
FT Modified-site 36 /note= "amidated residue"  
XX

PN US6114304-A.  
XX  
XX 05-SEP-2000.  
XX  
XX 07-SEP-1994; 94US-00302069.  
XX  
XX 07-SEP-1993; 93US-00118381.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX Young AA, Rink TJ, Brown KAK, Kolterman OG;  
XX  
XX WPI; 2000-601336/57.  
XX  
XX  
XX Treating gastrointestinal disorder e.g. spasm by reducing gastric  
XX motility or delaying gastric emptying, postprandial dumping syndrome or  
XX postprandial hyperglycemia, by administering amylin or amylin agonist.  
XX  
XX Disclosure; Col 41-42; 50pp; English.  
XX  
XX The present sequence represents an amylin agonist analogue compound.  
XX Amylin or amylin agonists are administered for reducing gastric motility  
XX or delaying gastric emptying, and for treating postprandial dumping  
XX syndrome or postprandial hyperglycemia, by inducing amylin activity, in a  
XX mammal. The peptides are used to reduce gastric motility or for delaying  
XX gastric emptying in a mammal undergoing gastrointestinal diagnostic  
XX procedures, such as radiological examination or magnetic resonance  
XX imaging. They are also used for reducing gastric motility in  
XX gastrointestinal disorder, especially spasm, which is associated with a  
XX disorder of acute diverticulitis or disorders of biliary tract or  
XX sphincter of oddi. They are also used to treat postprandial dumping  
XX syndrome or postprandial hyperglycemia  
XX  
XX Sequence 36 AA;  
SQ  
Query Match 97.5%; Score 197; DB 3; Length 36;  
Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPTNVGSNTY 36  
RESULT 42  
ABB05495  
ID ABB05495 standard; peptide; 36 AA.  
XX  
XX ABB05495;  
AC  
XX  
XX 19-APR-2002 (first entry)  
DT  
XX  
XX Human amylin agonist des-Ilys25,28,29Pro-h-amylin.  
DE  
XX  
XX Human; amylin agonist; amylin peptide analogue; acetate; carbohydrate;  
XX liquid pharmaceutical formulation; polyhydric alcohol; phosphate;  
XX citrate; glutamate; buffer; antidiabetic; type II diabetes.  
XX  
XX Homo sapiens.  
OS  
XX Synthetic.  
XX  
XX US2001043934-A1.  
XX  
XX 22-NOV-2001.  
XX  
XX 09-JAN-1998; 98US-00005262.  
XX  
XX 08-JAN-1997; 97US-0035140P.  
XX  
XX (LITA/) L'ITALIEN J.  
XX (MUSU/) MUSUNURI S.  
XX (RUBY/) RUBY K.  
XX

PI L'italien J, Musunuri S, Ruby K;  
XX  
XX WPI; 2002-163554/21.  
XX  
XX New pharmaceutical formulation useful for treating patients with type II  
XX diabetes comprises amylin agonist, carbohydrate or polyhydric alcohol and  
XX a buffer.  
XX  
XX Disclosure; Page; 19pp; English.  
XX  
XX The present invention describes a liquid pharmaceutical formulation (A)  
XX comprising (wt/vol%) an amylin agonist (0.01 - 0.5), a carbohydrate or a  
XX polyhydric alcohol (1 - 10) and an acetate, phosphate, citrate or  
XX glutamate buffer (0.02 - 0.5) having a pH of 3 - 6. Also described is a  
XX commercial package containing the liquid pharmaceutical formulation (A).  
XX The package comprises a borosilicate glass vial having an open end, a  
XX stopper for multiseal compatible with the amylin and/or amylin agonist  
XX fixed in the open end of the vial and an aluminum band to retain the  
XX stopper in the far end of the vial. The package also comprises a  
XX cartridge for use in a pen injector. (A) has antidiabetic activity and  
XX can be used in the treatment of patients with type II diabetes. The  
XX formulation comprises amylin agonist which is biologically active, has a  
XX reduced tendency to form aggregates in water or at a pressure of greater  
XX than 2 psi and has a reduced tendency to precipitate in the presence of  
XX NaCl compared to human amylin. The formulation maintains stability upon  
XX storage under refrigerated or room-temperature conditions. The  
XX formulation retains short-term mixing compatibility with insulin and  
XX results in improved stability of the hormone and the patients no longer  
XX need to refrigerate the vial of insulin in use. The present sequence  
XX represents a human amylin peptide analogue, which can be used as an  
XX amylin agonist in the present invention. N.B. The present sequence is not  
XX given in the present specification but is derived from the 37 amino acid  
XX human amylin as stated in the invention  
XX  
XX Sequence 36 AA;  
SQ  
Query Match 97.5%; Score 197; DB 5; Length 36;  
Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 CNTATCATQRLANFLVHSSNNFGPILPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPTNVGSNTY 36  
RESULT 43  
AD136180  
ID AD136180 standard; peptide; 36 AA.  
XX  
XX AD136180;  
AC  
XX  
XX 15-APR-2004 (first entry)  
DT  
XX  
XX Human amylin agonist analogue #9.  
DE  
XX  
XX Gastric motility; gastric emptying; amylin agonist; therapy; diagnosis;  
XX antidiabetic; hypoglycemia; human.  
XX  
XX Homo sapiens.  
OS  
XX  
XX Key Location/Qualifiers  
XX FH Disulfide-bond 1..6  
XX FT Modified-site 36  
XX FT /note= "Amidated tyrosine"  
XX  
XX US6608029-B1.  
XX  
XX 19-AUG-2003.  
XX  
XX 22-MAY-2000; 2000US-00576062.  
XX  
XX 07-SEP-1993; 93US-00118381.  
XX  
XX 07-SEP-1994; 94US-00302069.  
XX  
XX



XX (AMYL-) AMYLIN PHARM INC.  
 PA Kolterman OG, Young AA, Rink TJ, Keating Brown KA;  
 XX WPI; 2004-118064/12.  
 XX  
 PT Reducing gastric motility or delaying gastric emptying in a mammal,  
 PT useful for treating post-prandial hyperglycemia, comprises administering  
 an amylin or an amylin agonist.  
 XX  
 PS Disclosure; SEQ ID NO 10; 51bp; English.  
 XX  
 CC The present invention is directed to novel methods for reducing gastric  
 CC motility and delaying gastric emptying, comprising the administration of  
 CC an amylin or an amylin agonist. The invention is useful for reducing  
 CC gastric motility and delaying gastric emptying for therapeutic and  
 CC diagnostic purposes. The invention is also useful for treating conditions  
 CC associated with elevated, inappropriate or undesired post-prandial blood  
 CC glucose levels and treating ingestion of a toxin. The present sequence is  
 CC human amylin agonist analogue.  
 XX  
 SQ Sequence 36 AA;  
 XX  
 Query Match 97.5%; Score 197; DB 8; Length 36;  
 Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 2 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37  
 Db 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 36  
 XX  
 RESULT 44  
 ADO51024  
 ID ADO51024 standard; peptide; 36 AA.  
 XX  
 AC ADO51024;  
 XX  
 DT 18-NOV-2004 (first entry)  
 XX  
 DE Human amylin agonist peptide analogue #10.  
 XX  
 KW Gastric motility; delay gastric emptying; amylin; agonist;  
 KW postprandial dumping syndrome; postprandial hyperglycemia;  
 KW gastrointestinal disorder; spasm; radiological examination;  
 KW magnetic resonance imaging; diabetes; therapy; human.  
 XX  
 OS Homo sapiens.  
 XX  
 FH Key Location/Qualifiers  
 FT Modified-site 36 /note= "C-terminal amide"  
 XX  
 PN US2004097415-A1.  
 XX  
 PD 20-MAY-2004.  
 XX  
 PF 18-AUG-2003; 2003US-00643681.  
 XX  
 PR 07-SEP-1993; 93US-00118381.  
 PR 07-SEP-1994; 94US-00302069.  
 PR 22-MAY-2000; 2000US-00576062.  
 XX  
 PA (KOLJ/) KOLTERMAN O G.  
 PA (YOUN/) YOUNG A A.  
 PA (RINK/) RINK T J.  
 PA (BROW/) KEATING BROWN K A.  
 XX  
 PI Kolterman OG, Young AA, Rink TJ, Keating Brown KA;  
 XX WPI; 2004-389180/36.  
 XX

PT Use of amylin agonist for reducing gastric motility, delaying gastric  
 PT emptying and for treating postprandial dumping syndrome and postprandial  
 PT hyperglycemia.  
 XX  
 XX Disclosure; SEQ ID NO 10; 35bp; English.  
 XX  
 CC The present invention is directed to novel methods for reducing gastric  
 CC motility and delaying gastric emptying which comprises the administration  
 CC of an amylin or an amylin agonist. The invention is useful for treating  
 CC postprandial dumping syndrome, postprandial hyperglycemia and reducing  
 CC gastric motility associated with gastrointestinal disorders such as spasm  
 CC or delaying gastric emptying in a mammal undergoing a gastrointestinal  
 CC diagnostic procedure such as radiological examination and magnetic  
 CC resonance imaging. The invention is also useful for lowering postprandial  
 CC blood glucose levels during treatment of diabetes. The present sequence  
 CC is human amylin agonist peptide analogue. This sequence is used in the  
 CC invention.  
 XX  
 SQ Sequence 36 AA;  
 XX  
 Query Match 97.5%; Score 197; DB 8; Length 36;  
 Best Local Similarity 100.0%; Pred. No. 3.2e-20;  
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 2 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37  
 Db 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 36  
 XX  
 RESULT 45  
 AAR37794  
 ID AAR37794 standard; peptide; 36 AA.  
 XX  
 AC AAR37794;  
 XX  
 DT 25-MAR-2003 (revised)  
 DT 07-SEP-1993 (first entry)  
 XX  
 DE (Des-Lys1), Pro25, Val26, Pro28, Pro29 human amylin analogue.  
 XX  
 KW Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;  
 KW hyperglycaemic agent.  
 XX  
 OS Homo sapiens.  
 XX  
 FH Key Location/Qualifiers  
 FT Disulfide-bond 1..6  
 FT Misc-difference 24 /note= "Pro replaces wild-type Ala"  
 FT Misc-difference 25 /note= "val replaces wild-type Ile"  
 FT Misc-difference 27 /note= "Pro replaces wild-type Ser"  
 FT Misc-difference 28 /note= "Pro replaces wild-type Ser"  
 FT Modified-site 36 /note= "amidated"  
 XX  
 PN WO9310146-A1.  
 XX  
 PD 27-MAY-1993.  
 XX  
 PR 19-NOV-1992; 92WO-US009842.  
 PR 19-NOV-1991; 91US-00794266.  
 XX  
 PA (AMYL-) AMYLIN PHARM INC.  
 XX  
 PI Gaeta LSL, Jones H, Albrecht E;  
 XX WPI; 1993-182488/22.  
 XX  
 XX New amylin agonist peptide(s) - used for treatment and prevention of

PT hypoglycaemia and diabetes mellitus.  
XX  
PS Example 16; Fig 1 and Page 23; 43pp; English.  
XX  
CC This peptide is an example of amylin agonists of the invention which can  
CC be used as hyperglycaemic. The peptide is an analogue of human amylin  
CC which mimics the effects of the wild-type hormone. Preferred peptides are  
CC used in admixture with insulin for the treatment of diabetes mellitus or  
CC with glucagon for the treatment of hypoglycaemic conditions. See AAR37779  
CC -R37795 and AAR38809-R38826. (Updated on 25-MAR-2003 to correct PN  
CC field.)  
XX  
SQ Sequence 36 AA;  
XX  
Query Match 97.0%; Score 196; DB 2; Length 36;  
Best Local Similarity 97.2%; Pred. No. 4.4e-20;  
Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
XX  
QY 2 CNTATCATQRLANFLVHSSNNFGPIIPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPIIPPTNVGSNTY 36  
XX  
RESULT 46  
AAM22579  
ID AAM22579 standard; peptide; 37 AA.  
XX  
AC AAM22579;  
XX  
DT 12-OCT-1997 (first entry)  
XX  
DE Amylin agonist/CKK agonist hybrid peptide.  
XX  
KW appetite regulation; amylin agonist; hybrid; CKK; cholecystokinin.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 37 /note="Tyr-NH2"  
XX  
XX  
XX MO9640196-A1.  
XX  
XX 19-DEC-1996.  
XX  
XX 06-JUN-1996; 96WO-US009937.  
XX  
XX 07-JUN-1995; 95US-00477727.  
XX  
XX (AMYL-) AMYLIN PHARM INC.  
XX  
XX Rink TU, Young AA, Beeley NR, Prickett KS;  
XX  
XX WPI; 1997-051885/05.  
XX  
XX  
XX Compositions comprising amylin and cholecystokinin agonists - useful for  
XX reducing food intake, suppressing appetite and controlling body weight.  
XX  
XX Claim 81; Page 61; 66pp; English.  
XX  
XX The patent discloses a composition which can be used to reduce or  
XX suppress food intake, control appetite or control body weight in a  
XX mammal, comprising an amylin agonist and a cholecystokinin (CKK) agonist  
XX admixed in a form suitable for therapeutic administration. Also disclosed  
XX are new hybrid peptides comprising an amylin agonist peptide joined to a  
XX CKK agonist via a covalent linking group (see AAM22540 - AAM22551).  
XX Further disclosed are new hybrid peptides which incorporate features of  
XX amylin agonist peptides and CKK agonist peptides but which do not employ  
XX a linking group. These peptides are of general formula R1-C-R2-C-R3-R4-  
XX R5, in which R1 = a free N-terminus or an amidated N-terminus or amidated  
XX lysine (L), where amidation is with acetamide, propionamide, butyramide,  
XX isobutyramide or isocaproamide; R2 = an amino acid sequence selected from  
XX NTAT, GTAT, NVT, NMAAT, SNLST, ASLST and GNLST; R3 = an amino acid

CC sequence selected from ATQRLANFLVH and VLKLSQELHK; R4 = an amino acid  
CC sequence selected from SSNNFGPIIP and LOTYPR; and R5 = an amino acid  
CC sequence selected from DYNWMDNF-NH2, TNGWMDNF-NH2, TNGWMDNF-NH2, TNGWMDNF  
CC -NH2, TNGWMDNF-NH2, TNGWMDNF-NH2, TNGWMDNF-NH2, TNGWMDNF-NH2 and TNGWMDNF  
CC -NH2. Administration of amylin and CKK agonists in conjunction produces a  
CC greater effect than either administered alone; e.g. 0.1 microgram/kg of  
CC each peptide causes a substantial reduction of food intake about  
CC equivalent to that seen with 100 microgram/kg of either peptide alone.  
CC The present sequence represents one of 10 specific examples of hybrid  
CC peptides falling within the scope of the above formula  
XX  
SQ Sequence 37 AA;  
XX  
Query Match 97.0%; Score 196; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 4.5e-20;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
XX  
QY 1 KNTATCATQRLANFLVHSSNNFGPIIPPTNVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNNFGPIIPPTNVGSNDY 37  
XX  
RESULT 47  
ADH22050  
ID ADH22050 standard; peptide; 36 AA.  
XX  
AC ADH22050;  
XX  
DT 11-MAR-2004 (first entry)  
XX  
DE Amylin analogue pramlintide, SEQ ID NO:847.  
XX  
KW Fusion protein; human serum albumin; HSA; therapeutic protein;  
KW shelf-life; in vitro biological activity; in vivo biological activity;  
KW metabolic disorder; endocrine disorder; diabetes; type 1; type 2;  
KW diabetes-related condition; hyperglycaemia; neural disorder; neuropathy;  
KW retinopathy; cardiovascular disorder; heart disease; renal disorder;  
KW obesity; glucose level maintenance; weight loss; antidiabetic; cardiac;  
KW anorectic; ophthalmological; gene therapy.  
XX  
XX Homo sapiens.  
XX  
XX WO2003059934-A2.  
XX  
XX 24-JUL-2003.  
XX  
XX 23-DEC-2002; 2002WO-US040892.  
XX  
XX 21-DEC-2001; 2001US-0341811P.  
XX  
XX 26-JAN-2002; 2002US-0350358P.  
XX  
XX 26-FEB-2002; 2002US-0359370P.  
XX  
XX 28-FEB-2002; 2002US-0360000P.  
XX  
XX 27-MAR-2002; 2002US-0367500P.  
XX  
XX 08-APR-2002; 2002US-0370227P.  
XX  
XX 10-MAY-2002; 2002US-0378950P.  
XX  
XX 24-JUL-2002; 2002US-0398008P.  
XX  
XX 09-AUG-2002; 2002US-0402131P.  
XX  
XX 13-AUG-2002; 2002US-0402708P.  
XX  
XX 18-SEP-2002; 2002US-0411355P.  
XX  
XX 02-OCT-2002; 2002US-0414984P.  
XX  
XX 11-OCT-2002; 2002US-0417611P.  
XX  
XX 23-OCT-2002; 2002US-0420246P.  
XX  
XX 05-NOV-2002; 2002US-0423623P.  
XX  
XX (HMDA-) HUMAN GENOME SCI INC.  
XX  
XX Rosen CA, Haseltine WA;  
XX  
XX WPI; 2003-598501/56.  
XX  
XX New albumin fusion protein, useful for preparing a composition for  
XX treating diabetes mellitus.

PS Disclosure; SEQ ID NO 847; 1086pp; English.

The invention relates to fusion proteins comprising human serum albumin (ADH21530) and a therapeutic polypeptide such as a therapeutic protein, antibody or peptide or their variants or fragments. The therapeutic protein may be fused to the N-terminus, the C-terminus or both termini of albumin via a linker. The albumin component of the fusion proteins prolongs the shelf-life and the in vitro and vivo biological activity of the proteins compared with those of the corresponding therapeutic proteins on their own. The invention also relates to nucleic acids encoding albumin fusion proteins, vectors and host cells comprising an albumin fusion protein nucleic acid, compositions and kits comprising an albumin fusion protein, the method of extending the shelf-life of a therapeutic protein by fusion with albumin, and the treatment of disease using an albumin fusion protein. The albumin fusion proteins may be used in the treatment of metabolic/endocrine disorders, diabetes and diabetes related conditions. Specifically the albumin fusion proteins may be used to treat type 1 and type 2 diabetes, hyperglycaemia, neural disorders (especially neuropathy), retinopathy, cardiovascular disorders (especially heart disease), renal disorders and obesity. The proteins may also be used in a method of maintaining a basal glucose level in a patient and in a method for losing weight. The present sequence is related to the invention.

**SQ** Sequence 36 AA;

Query Match	96.5%	Score 195;	DB 7;	Length 36;
Best Local Similarity	100.0%	Pred. No. 6.1e-20;		
Matches 36;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0

Qy	Db
1 KCNTATCATQRLANFLVHSSNNFGPIIPTVGSNT 36	1 KCNTATCATQRLANFLVHSSNNFGPIIPTVGSNT 36

RESULT 48  
AAR29205  
ID AAR29205 standard; protein; 37 AA

AC	AAAR29205;
XX	
DT	25-MAR-2003 (revised)
DT	20-APR-1993 (first entry)
XX	
DE	Leu (23) Pro (25) Val (26) Pro (28, 29) -h-amylin for treating anorexia

KW Anorexia; cachexia; adipose; amylin.

**Homo sapiens.**

PN WO9220367-A1.

PD 26-NOV-1992.

PF 23-MAY-1992; 92WO-US004357

24-MAY-1991: 91US-00704995

XX 03-APR-1992; 94US-00862500.

PA (AMYL-) AMYLIN PHARM INC.

PI Rink TJ, Young AA;

DR WPI; 1992-415470/50.

PT Use of amylin and opt. insulin for treating anorexia - increases plasma  
PT amylin and/or insulin levels, also for treating cachexia conditions,  
PT adipose tissue deficiency etc.

CC Treatment of a patient with anorexia or related condition comprises  
CC admin. of amylin or an analogue in an amt. sufficient to increase the

CC amylin reveal in the plasma of the patient. The pref. amylin analogues are  
CC given in AAR29197-222. Treating a patient deficient in adipose tissue  
CC comprises admin. of amylin or an analogue and/or insulin in an ant. and  
CC ratio sufficient to increase adipose tissue. Typical dosage units contain  
CC 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-  
CC 2003 to correct PN field.)

Sequence 37 AA;

Query Match	96.5%	Score 195;	DB 2;	Length 37;
Best Local Similarity	94.6%	Pred. No. 6.3e-20;		
Matches 35; Conservative	1;	Mismatches 1;	Indels 0;	Gaps 0

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Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPTPNVGSNTY 37
        |||||
Db      1 KCNTATCATQRLANFLVHSSNNIGPVLPTPNVGSNTY 37
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RESULT 49  
AAR29211  
ID AAR29211 standard; protein; 37 AA

AC AAR29211;  
XX  
XX 25-MAR-2003 (revised)  
DT 20-APR-1993 (first entry)  
XX  
XX Tie(17)Leu(23)Pro(25,28,29)-h-amylin for treating anorexia

KW Anorexia; cachexia; adipose; amylin.

**Homo sapiens.**

PN WO9220367-A1.

PD 26-NOV-1992

PF 23-MAY-1992; 92WO-US004357.

PR 24-MAY-1991; 91US-00704995.

PR 03-APR-1992; 92US-00862500.

PA (AMYL-) AMYLIN PHARM INC.

PI Rink TJ, Young AA,

DR WPI; 1992-415470/50.

PT Use of amylin and opt. insulin for treating anorexia - increases plasma  
PT amylin and/or insulin levels, also for treating cachexia conditions,  
PT adipose tissue deficiency etc.

Treatment of a patient with anorexia or related condition comprises admin. of amylin or an analogue in an amt. sufficient to increase the amylin level in the plasma of the patient. The pref. amylin analogues are given in AAR9197-222. Treating a patient deficient in adipose tissue comprises admin. of amylin or an analogue and/or insulin in an amt. and ratio sufficient to increase adipose tissue. Typical dosage units contain 0.1-10 mg of amylin analogue and 0.1-1 mg of insulin. (Updated on 25-MAR-2003 to correct PN field.)

Sequence 37 AA;

Query Match	96.5%	Score 195;	DB 2;	Length 37;
Best Local Similarity	94.6%	Pred. No. 6.3e-20;		
Matches 35;	Conservative 1;	Mismatches 1;	Indels 0;	Gaps 0;

**Oy**      1 KCATATCATQRLANFLVHSSNNCPILPPTNVGSNTY 37  
         ||||| : |||||  
**Dd**      1 KCATATCATQRLANFLIHSSNNICPILPPTNVGSNTY 37

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RESULT 50
AAR38809
ID AAR38809 standard; peptide; 37 AA.
XX
AC AAR38809;
XX
DT 25-MAR-2003 (revised)
DT 07-SEP-1993 (first entry)
XX
DE Leu23, Pro25, Val26, Pro28, Pro29 human amylin analogue.
XX
KM Hypoglycaemia; insulin; pancreatic amyloid; diabetes mellitus; glucagon;
KM hyperglycaemic agent.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Disulfide-bond 2..7
FT Misc-difference 23
FT Misc-difference 25 /note= "Leu replaces wild-type Phe"
FT Misc-difference 28 /note= "Pro replaces wild-type Ala"
FT Misc-difference 29 /note= "Pro replaces wild-type Ser"
FT Modified-site 37 /note= "Pro replaces wild-type Ser"
FT /note= "amidated"
XX
FN WO9310146-A1.
XX
PD 27-MAY-1993.
XX
PF 19-NOV-1992; 92MO-US009842.
XX
PR 19-NOV-1991; 91US-00794266.
XX
PA (AMYL-) AMYLIN PHARM INC.
XX
PI Gaeta LSL, Jones H, Albrecht E;
XX
DR WPI, 1993-18248/22.
XX
PT New amylin agonist peptide(s) - used for treatment and prevention of
PT hypoglycaemia and diabetes mellitus.
XX
PS Example 18; Fig 3 and Page 29; 43pp; English.
XX
CC This peptide is an example of amylin agonists of the invention which can
CC be used as hyperglycaemics. The peptide is an analogue of human amylin
CC which mimics the effects of the wild-type hormone. Preferred peptides are
CC used in admixture with insulin for the treatment of diabetes mellitus or
CC with glucagon for the treatment of hypoglycaemic conditions. See AAR37779
CC -R37795 and AAR38809-R38826. (Updated on 25-MAR-2003 to correct PN
CC field.)
XX
SQ Sequence 37 AA;

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Query Match 96.5%; Score 195; DB 2; Length 37;
Best Local Similarity 94.6%; Pred. No. 6.3e-20;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
OY 1 KCNTATCATQRLANFLVHSSNNGPILPTNVGSNTY 37
DB 1 KCNTATCATQRLANFLVHSSNNGPILPTNVGSNTY 37

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Search completed: May 12, 2006, 15:18:09  
Job time : 188 secs

GenCore version 5.1.8  
Copyright (c) 1993 - 2006 Bioacceleration Ltd.

OM protein - protein search, using sw model

Run on: May 12, 2006, 15:33:32 ; Search time 164 Seconds  
(without alignments)  
94.266 Million cell updates/sec

Title: US-08-870-762b-1  
Perfect score: 202  
Sequence: 1 KCNTATCATGATRLANFLVHSSNNPGFILPTWGSNTY 37

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 100 summaries

Database : Published Applications AA Main:\*  
1: /cgn2\_6/prodata/1/pubppaa/US07\_PUBCOMB.pep:\*  
2: /cgn2\_6/prodata/1/pubppaa/US08\_PUBCOMB.pep:\*  
3: /cgn2\_6/prodata/1/pubppaa/US09\_PUBCOMB.pep:\*  
4: /cgn2\_6/prodata/1/pubppaa/US10A\_PUBCOMB.pep:\*  
5: /cgn2\_6/prodata/1/pubppaa/US10B\_PUBCOMB.pep:\*  
6: /cgn2\_6/prodata/1/pubppaa/US11\_PUBCOMB.pep:\*  
  
Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	202	100.0	37	2	US-08-851-965-1 Sequence 1, Appl1
2	202	100.0	37	2	US-08-870-762A-1 Sequence 9, Appl1
3	202	100.0	37	3	US-09-454-533-9 Sequence 1, Appl1
4	202	100.0	37	4	US-10-649-138-9 Sequence 9, Appl1
5	202	100.0	37	4	US-10-643-681-1 Sequence 1, Appl1
6	202	100.0	37	5	US-10-991-597-14 Sequence 14, Appl1
7	202	100.0	37	5	US-10-991-597-47 Sequence 47, Appl1
8	202	100.0	37	5	US-10-775-204-2200 Sequence 2200, Ap
9	201	99.5	37	2	US-08-851-965-6 Sequence 6, Appl1
10	201	99.5	37	2	US-08-851-965-17 Sequence 17, Appl1
11	201	99.5	37	2	US-08-870-762A-5 Sequence 5, Appl1
12	201	99.5	37	3	US-09-454-533-12 Sequence 12, Appl1
13	201	99.5	37	4	US-10-649-138-12 Sequence 12, Appl1
14	201	99.5	37	4	US-10-643-681-7 Sequence 7, Appl1
15	201	99.5	37	4	US-10-649-138-16 Sequence 18, Appl1
16	201	99.5	37	5	US-10-991-597-17 Sequence 17, Appl1
17	201	97.5	36	2	US-08-851-965-9 Sequence 9, Appl1
18	197	97.5	36	2	US-08-870-762A-15 Sequence 15, Appl1
19	197	97.5	36	4	US-10-643-681-10 Sequence 10, Appl1
20	195	96.5	36	5	US-10-775-180-847 Sequence 847, App
21	195	96.5	37	2	US-08-851-965-10 Sequence 10, Appl1
22	195	96.5	37	2	US-08-851-965-16 Sequence 16, Appl1
23	195	96.5	37	3	US-09-454-533-16 Sequence 16, Appl1
24	195	96.5	37	3	US-09-454-533-22 Sequence 22, Appl1
25	195	96.5	37	4	US-10-649-138-16 Sequence 16, Appl1
26	195	96.5	37	4	US-10-649-138-22 Sequence 22, Appl1
27	195	96.5	37	4	US-10-643-681-11 Sequence 11, Appl1

28	195	96.5	37	4	US-10-643-681-17 Sequence 17, Appl1
29	195	96.5	37	5	US-10-850-055-34 Sequence 34, Appl1
30	195	96.5	37	5	US-10-991-597-21 Sequence 21, Appl1
31	195	96.5	37	5	US-10-991-597-27 Sequence 27, Appl1
32	195	96.5	37	5	US-10-993-667-34 Sequence 34, Appl1
33	194	96.0	37	2	US-08-851-965-7 Sequence 7, Appl1
34	194	96.0	37	2	US-08-851-965-34 Sequence 34, Appl1
35	194	96.0	37	2	US-08-870-762A-2 Sequence 2, Appl1
36	194	96.0	37	2	US-08-870-762A-11 Sequence 11, Appl1
37	194	96.0	37	3	US-09-454-533-13 Sequence 13, Appl1
38	194	96.0	37	3	US-09-454-533-38 Sequence 38, Appl1
39	194	96.0	37	4	US-10-649-138-13 Sequence 13, Appl1
40	194	96.0	37	4	US-10-649-138-38 Sequence 38, Appl1
41	194	96.0	37	4	US-10-643-681-8 Sequence 8, Appl1
42	194	96.0	37	5	US-10-991-597-18 Sequence 18, Appl1
43	194	96.0	37	5	US-10-991-597-43 Sequence 43, Appl1
44	193	95.5	36	3	US-09-454-533-15 Sequence 15, Appl1
45	193	95.5	36	3	US-10-649-138-15 Sequence 15, Appl1
46	193	95.5	36	5	US-10-991-597-20 Sequence 20, Appl1
47	192	95.0	36	3	US-09-454-533-40 Sequence 40, Appl1
48	192	95.0	36	4	US-10-649-138-40 Sequence 40, Appl1
49	192	95.0	36	5	US-10-991-597-45 Sequence 45, Appl1
50	191	94.6	37	3	US-09-454-533-23 Sequence 23, Appl1
51	191	94.6	37	4	US-10-649-138-23 Sequence 23, Appl1
52	191	94.6	37	5	US-10-991-597-28 Sequence 28, Appl1
53	190	94.1	36	2	US-08-851-965-18 Sequence 18, Appl1
54	190	94.1	36	2	US-09-454-533-24 Sequence 24, Appl1
55	190	94.1	36	4	US-10-649-138-24 Sequence 24, Appl1
56	190	94.1	36	4	US-10-643-681-11 Sequence 11, Appl1
57	190	94.1	36	5	US-10-991-597-29 Sequence 29, Appl1
58	189	93.6	36	2	US-08-851-965-35 Sequence 35, Appl1
59	189	93.6	36	2	US-08-870-762A-12 Sequence 12, Appl1
60	189	93.6	36	2	US-08-870-762A-14 Sequence 14, Appl1
61	189	93.6	36	2	US-09-454-533-39 Sequence 39, Appl1
62	189	93.6	36	4	US-10-649-138-39 Sequence 39, Appl1
63	189	93.6	36	4	US-10-643-681-9 Sequence 9, Appl1
64	189	93.6	36	5	US-10-991-597-44 Sequence 44, Appl1
65	188	93.1	37	2	US-08-851-965-14 Sequence 14, Appl1
66	188	93.1	37	3	US-09-454-533-20 Sequence 20, Appl1
67	188	93.1	37	4	US-10-649-138-20 Sequence 20, Appl1
68	188	93.1	37	4	US-10-643-681-15 Sequence 15, Appl1
69	188	93.1	37	4	US-10-643-681-15 Sequence 15, Appl1
70	188	93.1	37	5	US-10-991-597-25 Sequence 25, Appl1
71	187	92.6	37	2	US-08-851-965-11 Sequence 11, Appl1
72	187	92.6	37	3	US-09-813-345-17 Sequence 17, Appl1
73	187	92.6	37	3	US-09-454-533-4 Sequence 4, Appl1
74	187	92.6	37	3	US-09-454-533-17 Sequence 17, Appl1
75	187	92.6	37	4	US-10-306-645A-1 Sequence 4, Appl1
76	187	92.6	37	4	US-10-649-138-4 Sequence 4, Appl1
77	187	92.6	37	4	US-10-649-138-17 Sequence 17, Appl1
78	187	92.6	37	4	US-10-643-681-12 Sequence 12, Appl1
79	187	92.6	37	5	US-10-991-597-9 Sequence 9, Appl1
80	187	92.6	37	5	US-10-991-597-22 Sequence 22, Appl1
81	187	92.6	37	6	US-11-066-697-321 Sequence 321, App
82	187	92.6	37	6	US-11-066-697-334 Sequence 334, App
83	186	92.1	93	5	US-10-481-666-5 Sequence 5, Appl1
84	186	92.1	37	2	US-08-851-965-3 Sequence 3, Appl1
85	186	92.1	37	2	US-08-851-965-4 Sequence 4, Appl1
86	186	92.1	37	2	US-08-851-965-21 Sequence 21, Appl1
87	186	92.1	37	2	US-08-851-965-33 Sequence 33, Appl1
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89	186	92.1	37	2	US-08-870-762A-4 Sequence 4, Appl1
90	186	92.1	37	2	US-08-870-762A-10 Sequence 10, Appl1
91	186	92.1	37	3	US-09-454-533-8 Sequence 8, Appl1
92	186	92.1	37	3	US-09-454-533-10 Sequence 10, Appl1
93	186	92.1	37	3	US-09-454-533-27 Sequence 27, Appl1
94	186	92.1	37	3	US-09-454-533-37 Sequence 37, Appl1
95	186	92.1	37	4	US-10-649-138-8 Sequence 8, Appl1
96	186	92.1	37	4	US-10-649-138-10 Sequence 10, Appl1
97	186	92.1	37	4	US-10-649-138-27 Sequence 27, Appl1
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99	186	92.1	37	4	US-10-643-681-3 Sequence 3, Appl1
100	186	92.1	37	4	US-10-643-681-5 Sequence 5, Appl1

## ALIGNMENTS

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RESULT 1
US-08-851-965-1
; Sequence 1, Application US/08851965
; Publication No. US2002010133A1
; GENERAL INFORMATION:
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: GEDULIN, Bronislava
; APPLICANT: BEYRON, Gareth Wyn
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: USING AMYLIN OR AMYLIN
; TITLE OF INVENTION: AGONISTS
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/851,965
; FILING DATE: 06-MAY-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 224/042
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 619/552-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; US-08-851-965-1

Query Match
Best Local Similarity 100.0%; Score 202; DB 2; Length 37;
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37

RESULT 2
US-08-870-762A-1
; Sequence 1, Application US/08870762A
; Publication No. US20030026812A1
; GENERAL INFORMATION:
; APPLICANT: Duft, Bradford
; APPLICANT: Koltzman, Orville
; TITLE OF INVENTION: METHODS FOR TREATING OBESITY
```

```
; NUMBER OF SEQUENCES: 15
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/870,762A
; FILING DATE: 06-JUN-1997
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 226/104
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-2200
; TELEFAX: 619-552-0159
; TELEX:
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; US-08-870-762A-1

Query Match
Best Local Similarity 100.0%; Score 202; DB 2; Length 37;
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVNGSNTY 37

RESULT 3
US-09-454-533-9
; Sequence 9, Application US/09454533
; Publication No. US20020187923A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USBS THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
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CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 9:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
US-09-454-533-9

Query Match 100.0%; Score 202; DB 3; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 4  
US-10-649-138-9  
Sequence 9, Application US/10649138  
Publication No. US20040038900A1  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/649,138  
FILING DATE: 26-Aug-2003  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991

APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 9:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
US-10-649-138-9

Query Match 100.0%; Score 202; DB 4; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 5  
US-10-643-681-1  
Sequence 1, Application US/10643681  
Publication No. US20040097415A1  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
YOUNG, Andrew A.  
RINK, Timothy J.  
BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/643,681  
FILING DATE: 18-Aug-2003  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

; INFORMATION FOR SEQ ID NO: 1:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
US-10-643-681-1

Query Match 100.0%; Score 202; DB 4; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 6  
US-10-991-597-14  
; Sequence 14, Application US/10991597  
; Publication No. US20050143303A1  
; GENERAL INFORMATION:  
; APPLICANT: Quay, Steven C.  
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF  
; FILE REFERENCE: 03-14US  
; CURRENT APPLICATION NUMBER: US/10/991,597  
; PRIOR FILING DATE: 2004-11-18  
; PRIOR APPLICATION NUMBER: 60/532,337  
; PRIOR FILING DATE: 2003-12-26  
; NUMBER OF SEQ ID NOS: 47  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 14  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-10-991-597-14

Query Match 100.0%; Score 202; DB 5; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 7  
US-10-991-597-47  
; Sequence 47, Application US/10991597  
; Publication No. US20050143303A1  
; GENERAL INFORMATION:  
; APPLICANT: Quay, Steven C.  
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF  
; FILE REFERENCE: 03-14US  
; CURRENT APPLICATION NUMBER: US/10/991,597  
; PRIOR FILING DATE: 2004-11-18  
; PRIOR APPLICATION NUMBER: 60/532,337  
; PRIOR FILING DATE: 2003-12-26  
; NUMBER OF SEQ ID NOS: 47  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 47  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens

US-10-991-597-47

Query Match 100.0%; Score 202; DB 5; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 8  
US-10-775-204-2200  
; Sequence 2200, Application US/10775204  
; Publication No. US20050186664A1  
; GENERAL INFORMATION:  
; APPLICANT: Rosen, Craig A.  
; APPLICANT: Haseltine, William A.  
; APPLICANT: Balance, David J.  
; APPLICANT: Turner, Andrew J.  
; TITLE OF INVENTION: Albumin Fusion Proteins  
; FILE REFERENCE: p564  
; CURRENT APPLICATION NUMBER: US/10/775,204  
; PRIOR FILING DATE: 2004-02-11  
; PRIOR APPLICATION NUMBER: 60/341,811  
; PRIOR FILING DATE: 2001-12-21  
; PRIOR APPLICATION NUMBER: 60/360,000  
; PRIOR FILING DATE: 2002-02-28  
; PRIOR APPLICATION NUMBER: 60/378,950  
; PRIOR FILING DATE: 2002-05-10  
; PRIOR APPLICATION NUMBER: 60/398,008  
; PRIOR FILING DATE: 2002-07-24  
; PRIOR APPLICATION NUMBER: 60/411,355  
; PRIOR FILING DATE: 2002-09-18  
; PRIOR APPLICATION NUMBER: 60/414,984  
; PRIOR FILING DATE: 2002-10-02  
; PRIOR APPLICATION NUMBER: 60/417,611  
; PRIOR FILING DATE: 2002-10-11  
; PRIOR APPLICATION NUMBER: 60/420,246  
; PRIOR FILING DATE: 2002-10-23  
; PRIOR APPLICATION NUMBER: 60/423,623  
; PRIOR FILING DATE: 2002-11-05  
; PRIOR APPLICATION NUMBER: 60/351,360  
; PRIOR FILING DATE: 2002-01-28  
; Remaining Prior Application data removed - See File Wrapper or PALM.  
; NUMBER OF SEQ ID NOS: 2222  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 2200  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-10-775-204-2200

Query Match 100.0%; Score 202; DB 5; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.5e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 9  
US-08-851-965-6  
; Sequence 6, Application US/08851965  
; Publication No. US20020010133A1  
; GENERAL INFORMATION:  
; APPLICANT: YOUNG, Andrew A.  
; APPLICANT: GEDULIN, Bronislava  
; APPLICANT: BEYNON, Gareth WYN  
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS  
; TITLE OF INVENTION: USING AMYLIN OR AMYLIN  
; AGONISTS



```

; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: FLOPPY disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/851,965
; FILING DATE: 06-MAY-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 224/042
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION:
; US-08-851-965-6

Query Match 99.5%; Score 201; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 2e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37
DB 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37

RESULT 10
US-08-851-965-17
; Sequence 17, Application US/08851965
; Publication No. US2002001013A1
; GENERAL INFORMATION:
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: GEDULIN, Bronislava
; APPLICANT: BEYNON, Gareth Wym
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: USING AMYLIN OR AMYLIN
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: FLOPPY disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
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; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/851,965
; FILING DATE: 06-MAY-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 224/042
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 17:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION:
; US-08-851-965-17

Query Match 99.5%; Score 201; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 2e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37
DB 1 KCNTATCATQRLANFLVHSSNNGPILPPTNGSNTY 37

RESULT 11
US-08-870-762A-5
; Sequence 5, Application US/08870762A
; Publication No. US20030026812A1
; GENERAL INFORMATION:
; APPLICANT: Duft, Bradford
; APPLICANT: Kolterman, Orville
; TITLE OF INVENTION: METHODS FOR TREATING OBESITY
; NUMBER OF SEQUENCES: 15
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FASTSEQ Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/870,762A
; FILING DATE: 06-JUN-1997
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 226/104
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-2200
; TELEFAX: 619-552-0159
; TELEX:
; INFORMATION FOR SEQ ID NO: 5:
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SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-870-762A-5

Query Match US-5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNFGPVLPTVGSNTY 37

RESULT 12  
US-09-454-533-12  
Sequence 12, Application US/09454533  
Publication No. US20020187923A1  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 12:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)

SEQUENCE DESCRIPTION: SEQ ID NO: 12:  
US-09-454-533-12

Query Match US-5%; Score 201; DB 3; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNFGPVLPTVGSNTY 37

RESULT 13  
US-10-649-138-12  
Sequence 12, Application US/10649138  
Publication No. US20040038900A1  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/649,138  
FILING DATE: 26-Aug-2003  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 12:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 12:  
US-10-649-138-12

Query Match US-5%; Score 201; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTVGSNTY 37  
|||||

Db 1 KCNTATCATQRLANFLVHSSNNFGPVLPTTVGSNTY 37

RESULT 14  
US-10-643-681-7  
; Sequence 7, Application US/10643681  
; Publication No. US20040097415A1  
; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; YOUNG, Andrew A.  
; RINK, Timothy J.  
; BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/10/643,681  
; FILING DATE: 18-Aug-2003  
; CLASSIFICATION: 514  
; PRIORITY APPLICATION DATA:  
; APPLICATION NUMBER: US/08/302,069A  
; FILING DATE: 07-SEP-1994  
; APPLICATION NUMBER: 08/118,381  
; FILING DATE: 07-SEP-1993  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 209/146  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 7:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:  
US-10-643-681-7

Query Match 99.5%; Score 201; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPVLPTTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPVLPTTVGSNTY 37

RESULT 15  
US-10-643-681-18  
; Sequence 18, Application US/10643681  
; Publication No. US20040097415A1  
; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; YOUNG, Andrew A.

; RINK, Timothy J.  
; BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/10/643,681  
; FILING DATE: 18-Aug-2003  
; CLASSIFICATION: 514  
; PRIORITY APPLICATION DATA:  
; APPLICATION NUMBER: US/08/302,069A  
; FILING DATE: 07-SEP-1994  
; APPLICATION NUMBER: 08/118,381  
; FILING DATE: 07-SEP-1993  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 209/146  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 18:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 18:  
US-10-643-681-18

Query Match 99.5%; Score 201; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPVLPTTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPVLPTTVGSNTY 37

RESULT 16  
US-10-991-597-17  
; Sequence 17, Application US/10991597  
; Publication No. US20050143303A1  
; GENERAL INFORMATION:  
; APPLICANT: Quay, Steven C.  
; APPLICANT: Costantino, Henry R.  
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF  
; FILE REFERENCE: 03-14US  
; CURRENT APPLICATION NUMBER: US/10/991,597  
; PRIOR FILING DATE: 2004-11-18  
; PRIOR APPLICATION NUMBER: 60/532,337  
; PRIOR FILING DATE: 2003-12-26  
; NUMBER OF SEQ ID NOS: 47  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 17

LENGTH: 37  
TYPE: PR  
ORGANISM: Homo sapiens  
US-10-991-597-17

Query Match 99.5%; Score 201; DB 5; Length 37;  
Best Local Similarity 97.3%; Pred. No. 2e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37  
Db 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37

RESULT 17  
US-08-851-965-9  
Sequence 9, Application US/08851965  
Publication No. US20020010133A1  
GENERAL INFORMATION:  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: GEDULIN, Bronislava  
APPLICANT: BEYRON, Gareth Wym  
TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS  
TITLE OF INVENTION: USING AMYLIN OR AMYLIN  
TITLE OF INVENTION: AGONISTS  
NUMBER OF SEQUENCES: 35  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/851,965  
FILING DATE: 06-MAY-1997  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 224/042  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 9:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 1,6  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 36  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-851-965-9

Query Match 97.5%; Score 197; DB 2; Length 36;  
Best Local Similarity 100.0%; Pred. No. 7e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37  
Db 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 36

RESULT 18  
US-08-870-762A-15  
Sequence 15, Application US/08870762A  
Publication No. US20030026812A1  
GENERAL INFORMATION:  
APPLICANT: Duft, Bradford  
APPLICANT: Koltzman, Orville  
TITLE OF INVENTION: METHODS FOR TREATING OBESITY  
NUMBER OF SEQUENCES: 15  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FASTSEQ Version 2.0  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/870,762A  
FILING DATE: 06-JUN-1997  
CLASSIFICATION: 424  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 226/104  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-2200  
TELEFAX: 619-552-0159  
TELEX:

INFORMATION FOR SEQ ID NO: 15:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 1,6  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 36  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-870-762A-15

Query Match 97.5%; Score 197; DB 2; Length 36;  
Best Local Similarity 100.0%; Pred. No. 7e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 37  
Db 1 CNTATCATORLANFLVHSSNFGPILPTNVGSNTY 36

RESULT 19  
US-10-643-681-10  
Sequence 10, Application US/10643681  
Publication No. US20040097415A1  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: BROWN, Kathleen Ann Keiting  
APPLICANT: RINK, Timothy J.  
TITLE OF INVENTION: METHODS FOR REGULATING  
GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30

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/
/ CORRESPONDENCE ADDRESS:
/ ADDRESSEE: LYON & LYON
/ STREET: 633 WEST FIFTH STREET
/ CITY: LOS ANGELES
/ STATE: CALIFORNIA
/ COUNTRY: USA
/ ZIP: 90017
/
/ COMPUTER READABLE FORM:
/ MEDIUM TYPE: Floppy disk
/ COMPUTER: IBM PC compatible
/ OPERATING SYSTEM: PC-DOS/MS-DOS
/ SOFTWARE: PatentIn Release #1.0, Version #1.25
/
/ CURRENT APPLICATION DATA:
/ APPLICATION NUMBER: US/10/643,681
/ FILING DATE: 18-Aug-2003
/ CLASSIFICATION: 514
/
/ PRIOR APPLICATION DATA:
/ APPLICATION NUMBER: US/08/302,069A
/ FILING DATE: 07-SEP-1994
/ APPLICATION NUMBER: 08/118,381
/ FILING DATE: 07-SEP-1993
/
/ ATTORNEY/AGENT INFORMATION:
/ NAME: DUFT, BRADFORD J.
/ REGISTRATION NUMBER: 32,219
/ REFERENCE/DOCKET NUMBER: 209/146
/ TELECOMMUNICATION INFORMATION:
/ TELEPHONE: 619/552-2200
/ TELEFAX: 213/955-0440
/ TELEX: 67-3510
/
/ INFORMATION FOR SEQ ID NO: 10:
/ SEQUENCE CHARACTERISTICS:
/ LENGTH: 36 amino acids
/ TYPE: amino acid
/ STRANDEDNESS: single
/ TOPOLOGY: linear
/ MOLECULE TYPE: peptide
/ FEATURE:
/ LOCATION: 36
/ OTHER INFORMATION: amidated Tyr (Tyrosinamide)
/
/ SEQUENCE DESCRIPTION: SEQ ID NO: 10:
/
/ US-10-643-681-10
/
/
/ Query Match 97.5%; Score 197; DB 4; Length 36;
/ Best Local Similarity 100.0%; Pred. No. 7e-20;
/ Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
/
/ QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37
/ DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 36
/
/ RESULT 20
/ US-10-775-180-847
/ Sequence 847, Application US/10775180
/ Publication No. US20050054570A1
/ GENERAL INFORMATION:
/ APPLICANT: Rosen, Craig A.
/ APPLICANT: Haseltine, William A.
/ TITLE OF INVENTION: Albumin Fusion Proteins
/ FILE REFERENCE: PF574
/ CURRENT APPLICATION NUMBER: US/10/775,180
/ CURRENT FILING DATE: 2004-02-11
/ PRIOR APPLICATION NUMBER: PCT/US02/40892
/ PRIOR FILING DATE: 2002-12-23
/ PRIOR APPLICATION NUMBER: 60/341,811
/ PRIOR FILING DATE: 2001-12-21
/ PRIOR APPLICATION NUMBER: 60/360,000
/ PRIOR FILING DATE: 2002-02-28
/ PRIOR APPLICATION NUMBER: 60/378,950
/ PRIOR FILING DATE: 2002-05-10
/ PRIOR APPLICATION NUMBER: 60/398,008
/ PRIOR FILING DATE: 2002-07-24
/ PRIOR APPLICATION NUMBER: 60/411,355
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/ PRIOR FILING DATE: 2002-09-18
/ PRIOR APPLICATION NUMBER: 60/414,984
/ PRIOR FILING DATE: 2002-10-02
/ PRIOR APPLICATION NUMBER: 60/417,611
/ PRIOR FILING DATE: 2002-10-11
/ PRIOR APPLICATION NUMBER: 60/420,246
/ PRIOR FILING DATE: 2002-10-23
/ PRIOR APPLICATION NUMBER: 60/423,623
/ PRIOR FILING DATE: 2002-11-05
/ Remaining Prior Application data removed - See file wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 858
/ SOFTWARE: PatentIn Ver. 2.0
/ SEQ ID NO 847
/ LENGTH: 36
/ TYPE: PRT
/ ORGANISM: Homo sapiens
/
/ US-10-775-180-847
/
/
/ Query Match 96.5%; Score 195; DB 5; Length 36;
/ Best Local Similarity 100.0%; Pred. No. 1.3e-19;
/ Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
/
/ QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNT 36
/ DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNT 36
/
/ RESULT 21
/ US-08-851-965-10
/ Sequence 10, Application US/08851965
/ Publication No. US20020010133A1
/ GENERAL INFORMATION:
/ APPLICANT: YOUNG, Andrew A.
/ APPLICANT: GEBULIN, Bronislava
/ APPLICANT: BEYRON, Gareth Wym
/ TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
/ TITLE OF INVENTION: USING AMYLIN OR AMYLIN
/ TITLE OF INVENTION: AGONISTS
/ NUMBER OF SEQUENCES: 35
/ CORRESPONDENCE ADDRESSES:
/ ADDRESSEE: LYON & LYON
/ STREET: 633 WEST FIFTH STREET
/ CITY: LOS ANGELES
/ STATE: CALIFORNIA
/ COUNTRY: USA
/ ZIP: 90017
/
/ COMPUTER READABLE FORM:
/ MEDIUM TYPE: Floppy disk
/ COMPUTER: IBM PC compatible
/ OPERATING SYSTEM: PC-DOS/MS-DOS
/ SOFTWARE: PatentIn Release #1.0, Version #1.25
/ CURRENT APPLICATION DATA:
/ APPLICATION NUMBER: US/08/851,965
/ FILING DATE: 06-MAY-1997
/ CLASSIFICATION: 514
/ ATTORNEY/AGENT INFORMATION:
/ NAME: DUFT, BRADFORD J.
/ REGISTRATION NUMBER: 32,219
/ REFERENCE/DOCKET NUMBER: 224/042
/ TELECOMMUNICATION INFORMATION:
/ TELEPHONE: 619/552-2200
/ TELEFAX: 213/955-0440
/ TELEX: 67-3510
/
/ INFORMATION FOR SEQ ID NO: 10:
/ SEQUENCE CHARACTERISTICS:
/ LENGTH: 37 amino acids
/ TYPE: amino acid
/ STRANDEDNESS: single
/ TOPOLOGY: linear
/ MOLECULE TYPE: procein
/ FEATURE:
/ LOCATION: 2,7
/ OTHER INFORMATION: disulfide bridge between
/
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OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-851-965-10

Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNNLGPVLPPTVGSNTY 37

RESULT 22

US-08-851-965-16  
Sequence 16, Application US/08851965  
Publication No. US20020010133A1

GENERAL INFORMATION:  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: GEDULIN, Bronislava  
APPLICANT: BEYRON, Gareth Wym  
TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS  
TITLE OF INVENTION: USING AMYLIN OR AMYLIN  
TITLE OF INVENTION: AGONISTS  
NUMBER OF SEQUENCES: 35  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/851,965  
FILING DATE: 06-MAY-1997  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 224/042  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440

TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 16:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-851-965-16

Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNNLGPVLPPTVGSNTY 37

RESULT 23  
US-09-454-533-16  
Sequence 16, Application US/09454533  
Publication No. US20020187923A1

GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR

NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440

TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 16:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 16:  
US-09-454-533-16

Query Match 96.5%; Score 195; DB 3; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNNLGPVLPPTVGSNTY 37

RESULT 24  
US-09-454-533-22

Sequence 22, Application US/09454533  
Publication No. US20020187923A1

GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:

ADDRESSER: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991

ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 22:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:

LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)

SEQUENCE DESCRIPTION: SEQ ID NO: 22:  
US-09-454-533-22

Query Match 96.5%; Score 195; DB 3; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 25  
US-10-649-138-16  
Sequence 16, Application US/10649138  
Publication No. US20040038900A1  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSER: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/649,138  
FILING DATE: 26-Aug-2003  
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991

ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 16:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:

LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)

SEQUENCE DESCRIPTION: SEQ ID NO: 16:  
US-10-649-138-16

Query Match 96.5%; Score 195; DB 4; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 26  
US-10-649-138-22  
Sequence 22, Application US/10649138  
Publication No. US20040038900A1  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USERS THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSER: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/649,138  
FILING DATE: 26-Aug-2003  
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>

APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 22:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 22:  
US-10-643-138-22

Query Match 96.5%; Score 195; DB 4; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 27  
US-10-643-681-11  
Sequence 11, Application US/10643681  
Publication No. US20040097415A1  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
YOUNG, Andrew A.  
RINK, Timothy J.  
BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/643,681  
FILING DATE: 18-AUG-2003  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200

TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 11:  
US-10-643-681-11

Query Match 96.5%; Score 195; DB 4; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.4e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTVGSNTY 37

RESULT 28  
US-10-643-681-17  
Sequence 17, Application US/10643681  
Publication No. US20040097415A1  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
YOUNG, Andrew A.  
RINK, Timothy J.  
BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/643,681  
FILING DATE: 18-AUG-2003  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 17:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:



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; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 17:
US-10-643-681-17

Query Match          96.5%; Score 195; DB 4; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.4e-19;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 29
US-10-850-055-34
; Sequence 34, Application US/10850055
; Publication No. US20050009742A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Erlandsson, Rikard
; APPLICANT: Friisen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Kristina
; APPLICANT: Haggbblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortessman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Ossolinak, Amina
; APPLICANT: Patrone, Cesare
; APPLICANT: Ronnholm, Harriet
; APPLICANT: Wikstrom, Lillian
; APPLICANT: Zachrisson, Olof
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
; FILE REFERENCE: 21882-517 CIP
; CURRENT APPLICATION NUMBER: US/10/850,055
; CURRENT FILING DATE: 2004-05-19
; PRIOR APPLICATION NUMBER: US 10/718,071
; PRIOR FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: US 60/427,912
; PRIOR FILING DATE: 2002-11-20
; NUMBER OF SEQ ID NOS: 66
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 34
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-850-055-34

Query Match          96.5%; Score 195; DB 5; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.4e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 30
US-10-991-597-21
; Sequence 21, Application US/10991597
; Publication No. US20050143303A1
; GENERAL INFORMATION:
; APPLICANT: Quay, Steven C.
; APPLICANT: Costantino, Henry R.
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
; FILE REFERENCE: 03-14US
```

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; CURRENT APPLICATION NUMBER: US/10/991,597
; CURRENT FILING DATE: 2004-11-18
; PRIOR APPLICATION NUMBER: 60/532,337
; PRIOR FILING DATE: 2003-12-26
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 21
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-991-597-21

Query Match          96.5%; Score 195; DB 5; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.4e-19;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37

RESULT 31
US-10-991-597-27
; Sequence 27, Application US/10991597
; Publication No. US20050143303A1
; GENERAL INFORMATION:
; APPLICANT: Quay, Steven C.
; APPLICANT: Costantino, Henry R.
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
; FILE REFERENCE: 03-14US
; CURRENT APPLICATION NUMBER: US/10/991,597
; CURRENT FILING DATE: 2004-11-18
; PRIOR APPLICATION NUMBER: 60/532,337
; PRIOR FILING DATE: 2003-12-26
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 27
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-991-597-27

Query Match          96.5%; Score 195; DB 5; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.4e-19;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 32
US-10-993-667-34
; Sequence 34, Application US/10993667
; Publication No. US20050209142A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Erlandsson, Rikard
; APPLICANT: Friisen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Nina
; APPLICANT: Haggbblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortessman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Ossolinak, Amina
; APPLICANT: Patrone, Cesare
```

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; APPLICANT: Ronnholm, Harriet
; APPLICANT: Wikstrom, Lillian
; APPLICANT: Zachrisson, Olof
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
; FILE REFERENCE: 21882-517 CIP2
; CURRENT APPLICATION NUMBER: US/10/993,667
; CURRENT FILING DATE: 2004-11-19
; PRIOR APPLICATION NUMBER: US 10/850,055
; PRIOR FILING DATE: 2003-05-19
; PRIOR APPLICATION NUMBER: US 10/718,071
; PRIOR FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: US 60/427,912
; PRIOR FILING DATE: 2002-11-20
; NUMBER OF SEQ ID NOS: 81
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 34
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-993-667-34

Query Match          96.5%; Score 195; DB 5; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.4e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 33
US-08-851-965-7
; Sequence 7, Application US/08851965
; Publication No. US20020010133A1
; GENERAL INFORMATION:
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: GEDULIN, Bronislava
; APPLICANT: BEYNON, Gareth Wyn
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: USING AMYLIN OR AMYLIN
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/851,965
; FILING DATE: 06-MAY-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 224/042
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide

Query Match          96.5%; Score 195; DB 5; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.4e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 34
US-08-851-965-34
; Sequence 34, Application US/08851965
; Publication No. US20020010133A1
; GENERAL INFORMATION:
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: GEDULIN, Bronislava
; APPLICANT: BEYNON, Gareth Wyn
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: METHOD FOR PREVENTING GASTRITIS
; TITLE OF INVENTION: USING AMYLIN OR AMYLIN
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/851,965
; FILING DATE: 06-MAY-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 224/042
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 34:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein

Query Match          96.0%; Score 194; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
US-08-851-965-7

Query Match          96.0%; Score 194; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
US-08-851-965-34

Query Match          96.0%; Score 194; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 35  
US-08-870-762A-2  
; Sequence 2, Application US/08870762A  
; Publication No. US20030026812A1  
; GENERAL INFORMATION:  
; APPLICANT: Duft, Bradford  
; APPLICANT: Kolterman, Orville  
; TITLE OF INVENTION: METHODS FOR TREATING OBESITY  
; NUMBER OF SEQUENCES: 15  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 90071-2066  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSEQ Version 2.0  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/870,762A  
; FILING DATE: 06-JUN-1997  
; CLASSIFICATION: 424  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER:  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 226/104  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619-552-2200  
; TELEFAX: 619-552-0159  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 2:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 2,7  
; OTHER INFORMATION: disulfide bridge between  
; OTHER INFORMATION: the Cys residues  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-870-762A-2

Query Match 96.0%; Score 194; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.9e-19;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 36  
US-08-870-762A-11  
; Sequence 11, Application US/08870762A  
; Publication No. US20030026812A1  
; GENERAL INFORMATION:  
; APPLICANT: Duft, Bradford  
; APPLICANT: Kolterman, Orville  
; TITLE OF INVENTION: METHODS FOR TREATING OBESITY  
; NUMBER OF SEQUENCES: 15

; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 90071-2066  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSEQ Version 2.0  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/870,762A  
; FILING DATE: 06-JUN-1997  
; CLASSIFICATION: 424  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER:  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 226/104  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619-552-2200  
; TELEFAX: 619-552-0159  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 11:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 2,7  
; OTHER INFORMATION: disulfide bridge between  
; OTHER INFORMATION: the Cys residues  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-870-762A-11

Query Match 96.0%; Score 194; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.9e-19;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 37  
US-09-454-533-13  
; Sequence 13, Application US/09454533  
; Publication No. US20020187923A1  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et Al.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; USES THEREFOR  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:

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; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-454-533-13

Query Match          96.0%; Score 194; DB 3; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 38
US-09-454-533-38
; Sequence 38, Application US/09454533
; Publication No. US20020187923A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-454-533-13

Query Match          96.0%; Score 194; DB 3; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 39
US-10-649-138-13
; Sequence 13, Application US/10649138
; Publication No. US20040038900A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/649,138
; FILING DATE: 26-Aug-2003
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
```

```
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 38
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-454-533-38

Query Match          96.0%; Score 194; DB 3; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 39
US-10-649-138-13
; Sequence 13, Application US/10649138
; Publication No. US20040038900A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/649,138
; FILING DATE: 26-Aug-2003
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
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;  
;  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
;  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
;  
; MOLECULE TYPE: peptide  
; FEATURE:  
;  
; LOCATION: 37  
;  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
;  
; SEQUENCE DESCRIPTION: SEQ ID NO: 13:  
US-10-649-138-13  
  
Query Match 96.0%; Score 194; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.9e-19;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
  
Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVRSSNFGPILPPTNVGSNTY 37  
|||||  
  
RESULT 40  
US-10-649-138-38  
; Sequence 38, Application US/10649138  
; Publication No. US20040038900A1  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et Al.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; USES THEREFOR  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/10/649,138  
; FILING DATE: 26-Aug-2003  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US/09/454,533  
; FILING DATE: 06-Dec-1999  
; APPLICATION NUMBER: 08/892,549  
; FILING DATE: <Unknown>  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-Nov-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-Mar-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 227/006  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 38  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 38:

US-10-649-138-38  
  
Query Match 96.0%; Score 194; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.9e-19;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
  
Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
  
RESULT 41  
US-10-643-681-8  
; Sequence 8, Application US/10643681  
; Publication No. US20040097415A1  
; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; YOUNG, Andrew A.  
; RINK, Timothy J.  
; BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/10/643,681  
; FILING DATE: 18-Aug-2003  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US/08/302,069A  
; FILING DATE: 07-SEP-1994  
; APPLICATION NUMBER: 08/118,381  
; FILING DATE: 07-SEP-1993  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 209/146  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 8:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 8:  
US-10-643-681-8  
  
Query Match 96.0%; Score 194; DB 4; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.9e-19;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
  
Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVRSSNFGPILPPTNVGSNTY 37  
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RESULT 42
US-10-991-597-18
; Sequence 18, Application US/10991597
; Publication No. US20050143303A1
; GENERAL INFORMATION:
; APPLICANT: Quay, Steven C.
; APPLICANT: Costantino, Henry R.
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
; TITLE OF INVENTION: GLUCOSE-REGULATING PEPTIDES
; FILE REFERENCE: 03-14US
; CURRENT APPLICATION NUMBER: US/10/991,597
; CURRENT FILING DATE: 2004-11-18
; PRIOR APPLICATION NUMBER: 60/532,337
; PRIOR FILING DATE: 2003-12-26
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 18
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-991-597-18

Query Match          96.0%; Score 194; DB 5; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 43
US-10-991-597-43
; Sequence 43, Application US/10991597
; Publication No. US20050143303A1
; GENERAL INFORMATION:
; APPLICANT: Quay, Steven C.
; APPLICANT: Costantino, Henry R.
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
; TITLE OF INVENTION: GLUCOSE-REGULATING PEPTIDES
; FILE REFERENCE: 03-14US
; CURRENT APPLICATION NUMBER: US/10/991,597
; CURRENT FILING DATE: 2004-11-18
; PRIOR APPLICATION NUMBER: 60/532,337
; PRIOR FILING DATE: 2003-12-26
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 43
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-991-597-43

Query Match          96.0%; Score 194; DB 5; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 44
US-09-454-533-15
; Sequence 15, Application US/09454533
; Publication No. US20020187923A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S. L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
US-09-454-533-15

Query Match          95.5%; Score 193; DB 3; Length 36;
Best Local Similarity 97.2%; Pred. No. 2.5e-19;
Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | |
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 45
US-10-649-138-15
; Sequence 15, Application US/10649138
; Publication No. US20040038900A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S. L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; APPLICANT: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: PatentIn Release #1.0, Version #1.25
US-10-649-138-15

Query Match          95.5%; Score 193; DB 3; Length 36;
Best Local Similarity 97.2%; Pred. No. 2.5e-19;
Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | |
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 46
US-09-454-533-15
; Sequence 15, Application US/10649138
; Publication No. US20040038900A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S. L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; APPLICANT: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: PatentIn Release #1.0, Version #1.25
US-09-454-533-15
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/ CURRENT APPLICATION DATA:
/ APPLICATION NUMBER: US/10/649,138
/ FILING DATE: 26-Aug-2003
/ CLASSIFICATION: <Unknown>
/ PRIOR APPLICATION DATA:
/ APPLICATION NUMBER: US/09/454,533
/ FILING DATE: 06-Dec-1999
/ APPLICATION NUMBER: 08/892,549
/ FILING DATE: <Unknown>
/ APPLICATION NUMBER: 07/794,266
/ FILING DATE: 19-Nov-1991
/ APPLICATION NUMBER: US 07/667,040
/ FILING DATE: 08-Mar-1991
/ ATTORNEY/AGENT INFORMATION:
/ NAME: DUFT, BRADFORD J.
/ REGISTRATION NUMBER: 32,219
/ REFERENCE/DOCKET NUMBER: 227/006
/ TELECOMMUNICATION INFORMATION:
/ TELEPHONE: 619/552-2200
/ TELEFAX: 213/955-0440
/ TELEX: 67-3510
/ INFORMATION FOR SEQ ID NO: 15:
/ SEQUENCE CHARACTERISTICS:
/ LENGTH: 36 amino acids
/ TYPE: amino acid
/ STRANDEDNESS: single
/ TOPOLOGY: linear
/ MOLECULE TYPE: protein
/ FEATURE:
/ LOCATION: 36
/ OTHER INFORMATION: amidated Tyr (Tyrosinamide)
/
/ US-10-649-138-15
/
/ Query Match 95.5%; Score 193; DB 4; Length 36;
/ Best Local Similarity 97.2%; Pred. No. 2.5e-19;
/ Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
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/ QY 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
/ |||||
/ DB 1 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 36
/ |||||
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/ RESULT 46
/ US-10-991-597-20
/ Sequence 20, Application US/10991597
/ Publication No. US20050143303A1
/ GENERAL INFORMATION:
/ APPLICANT: Quay, Steven C.
/ APPLICANT: Costantino, Henry R.
/ TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
/ FILE REFERENCE: 03-14US
/ CURRENT APPLICATION NUMBER: US/10/991,597
/ CURRENT FILING DATE: 2004-11-18
/ PRIOR APPLICATION NUMBER: 60/532,337
/ PRIOR FILING DATE: 2003-12-26
/ NUMBER OF SEQ ID NOS: 47
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 20
/ LENGTH: 36
/ TYPE: PRT
/ ORGANISM: Homo sapiens
/ US-10-991-597-20
/
/ Query Match 95.5%; Score 193; DB 5; Length 36;
/ Best Local Similarity 97.2%; Pred. No. 2.5e-19;
/ Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
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/ QY 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
/ |||||
/ DB 1 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 36
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RESULT 47
US-09-454-533-40
; Sequence 40, Application US/09454533
; Publication No. US20020187923A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA: US/09/454,533
; FILING DATE: 06-Dec-1999
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-Nov-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-Mar-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 40
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 36
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 40:
US-09-454-533-40
/
/ Query Match 95.0%; Score 192; DB 3; Length 36;
/ Best Local Similarity 94.4%; Pred. No. 3.5e-19;
/ Matches 34; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
/
/ QY 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
/ |||||
/ DB 1 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 36
/ |||||
/
/ RESULT 48
US-10-649-138-40
; Sequence 40, Application US/10649138
; Publication No. US20040038900A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
```

```
;
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
;
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
;
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/649,138
; FILING DATE: 26-Aug-2003
; CLASSIFICATION: <Unknown>
;
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
;
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
;
; INFORMATION FOR SEQ ID NO: 40
;
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 36
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 40:
US-10-649-138-40

Query Match          95.0%; Score 192; DB 4; Length 36;
Best Local Similarity 94.4%; Pred. No. 3.5e-19;
Matches 34; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 CNTATCATQRLANFLVHSSNNFGPVLPPSPNVGSNTY 36

RESULT 49
US-10-991-597-45
; Sequence 45, Application US/10991597
; Publication No. US20050143303A1
; GENERAL INFORMATION:
; APPLICANT: Quay, Steven C.
; APPLICANT: Costantino, Henry R.
; TITLE OF INVENTION: INTRANASAL ADMINISTRATION OF
; FILE REFERENCE: 03-14US
; CURRENT APPLICATION NUMBER: US/10/991,597
; CURRENT FILING DATE: 2004-11-18
; PRIOR APPLICATION NUMBER: 60/532,337
; PRIOR FILING DATE: 2003-12-26
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 45
; LENGTH: 36
; TYPE: PRT

; ORGANISM: Homo sapiens
US-10-991-597-45

Query Match          95.0%; Score 192; DB 5; Length 36;
Best Local Similarity 94.4%; Pred. No. 3.5e-19;
Matches 34; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 CNTATCATQRLANFLVHSSNNFGPVLPPSPNVGSNTY 36

RESULT 50
US-09-454-533-23
; Sequence 23, Application US/09454533
; Publication No. US20020187923A1
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
;
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/454,533
; FILING DATE: 06-Dec-1999
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/892,549
; FILING DATE: <Unknown>
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
;
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
;
; INFORMATION FOR SEQ ID NO: 23
;
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-09-454-533-23

Query Match          94.6%; Score 191; DB 3; Length 37;
Best Local Similarity 97.3%; Pred. No. 4.9e-19;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db      1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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Job time : 165 secs

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OM protein - protein search, using sw model

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(without alignments)  
64.337 Million cell updates/sec

Title: US-08-870-762B-1

Perfect score: 202

Sequence: 1 KCVATATCAQRLANFLVHSSNNFGPILPTTVGNSNTY 37

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Searched: 250354 seqs, 4694837 residues

Total number of hits satisfying chosen parameters: 250354

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 100 summaries

Database : Published Applications AA.New.\*

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- 2: /SIDSS/ptodata/2/pubpaa/US06\_NEW\_PUB.pep.\*
- 3: /SIDSS/ptodata/2/pubpaa/US07\_NEW\_PUB.pep.\*
- 4: /SIDSS/ptodata/2/pubpaa/US08\_NEW\_PUB.pep.\*
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- 6: /SIDSS/ptodata/2/pubpaa/US09\_NEW\_PUB.pep1.\*
- 7: /SIDSS/ptodata/2/pubpaa/US09\_NEW\_PUB.pep1.\*
- 8: /SIDSS/ptodata/2/pubpaa/US10\_NEW\_PUB.pep.\*
- 9: /SIDSS/ptodata/2/pubpaa/US10\_NEW\_PUB.pep1.\*
- 10: /SIDSS/ptodata/2/pubpaa/US11\_NEW\_PUB.pep1.\*
- 11: /SIDSS/ptodata/2/pubpaa/US11\_NEW\_PUB.pep1.\*
- 12: /SIDSS/ptodata/2/pubpaa/US60\_NEW\_PUB.pep1.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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3	201	99.5	37	10	US-11-055-093-75
4	201	99.5	37	10	US-11-055-093-92
5	197	97.5	36	10	US-11-055-093-74
6	195	96.5	37	10	US-11-055-093-85
7	195	96.5	37	10	US-11-055-093-91
8	195	96.5	37	11	US-11-288-495-34
9	194	96.0	37	10	US-11-055-093-72
10	194	96.0	37	10	US-11-055-093-83
11	190	94.1	36	10	US-11-055-093-93
12	189	93.6	36	10	US-11-055-093-73
13	189	93.6	36	10	US-11-055-093-84
14	188	93.1	37	10	US-11-055-093-89
15	187	92.6	37	10	US-11-055-093-44
16	187	92.6	37	10	US-11-055-093-86
17	186	92.1	37	10	US-11-055-093-70
18	186	92.1	37	10	US-11-055-093-82
19	186	92.1	37	10	US-11-055-093-96
20	186	92.1	37	10	US-11-055-093-189
21	182	90.1	36	10	US-11-055-093-87
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					Sequence 84, Appl
					Sequence 89, Appl
					Sequence 44, Appl
					Sequence 86, Appl
					Sequence 70, Appl
					Sequence 82, Appl
					Sequence 95, Appl
					Sequence 189, Appl
					Sequence 87, Appl

181	89.6	36	10	US-11-055-093-71	Sequence 71, Appl
180	89.1	37	10	US-11-055-093-90	Sequence 90, Appl
179	88.6	37	10	US-11-055-093-88	Sequence 88, Appl
178	88.1	37	9	US-10-516-768-26	Sequence 26, Appl
178	88.1	37	10	US-11-055-093-45	Sequence 45, Appl
178	88.1	37	11	US-11-033-105A-22	Sequence 22, Appl
178	88.1	37	11	US-11-288-495-7	Sequence 7, Appl
178	88.1	37	11	US-11-288-495-35	Sequence 35, Appl
178	88.1	89	11	US-11-145-463-93	Sequence 93, Appl
178	88.1	89	11	US-11-030-300-21	Sequence 21, Appl
178	88.1	103	11	US-11-145-463-95	Sequence 95, Appl
173	85.6	36	10	US-11-055-093-68	Sequence 68, Appl
173	85.6	37	9	US-10-760-085-8	Sequence 8, Appl
173	85.6	37	10	US-11-055-093-78	Sequence 78, Appl
173	85.6	37	10	US-11-055-093-81	Sequence 81, Appl
171	84.7	36	10	US-11-055-093-210	Sequence 210, Appl
170	84.2	37	10	US-11-055-093-95	Sequence 95, Appl
169	83.7	37	10	US-11-055-093-79	Sequence 79, Appl
169	83.7	37	10	US-11-055-093-102	Sequence 102, Appl
166	82.2	35	10	US-11-055-093-211	Sequence 211, Appl
166	82.2	37	10	US-11-055-093-101	Sequence 101, Appl
163	80.7	37	10	US-11-055-093-94	Sequence 94, Appl
162	80.2	37	10	US-11-055-093-76	Sequence 76, Appl
160	79.2	37	10	US-11-055-093-80	Sequence 80, Appl
159	78.7	37	10	US-11-055-093-97	Sequence 97, Appl
156	77.2	36	10	US-11-055-093-99	Sequence 99, Appl
153	75.7	37	10	US-11-055-093-98	Sequence 98, Appl
153	75.7	37	10	US-11-055-093-100	Sequence 100, Appl
113.5	56.2	43	10	US-11-055-093-37	Sequence 37, Appl
113.5	56.2	43	10	US-11-055-093-202	Sequence 202, Appl
106	52.5	20	10	US-11-055-093-212	Sequence 212, Appl
99	49.0	30	11	US-11-098-674-13	Sequence 13, Appl
98	48.5	18	10	US-11-055-093-213	Sequence 213, Appl
98	48.5	36	10	US-11-055-093-191	Sequence 191, Appl
91	45.0	37	10	US-11-055-093-112	Sequence 112, Appl
91	45.0	37	10	US-11-055-093-118	Sequence 118, Appl
90	44.6	17	10	US-11-055-093-214	Sequence 214, Appl
90	44.6	36	10	US-11-055-093-38	Sequence 38, Appl
87	43.1	37	10	US-11-055-093-111	Sequence 111, Appl
87	43.1	37	10	US-11-055-093-115	Sequence 115, Appl
86	42.6	16	10	US-11-055-093-215	Sequence 215, Appl
86	42.6	37	9	US-10-516-768-25	Sequence 25, Appl
86	42.6	37	9	US-10-760-085-39	Sequence 39, Appl
86	42.6	37	10	US-11-055-093-50	Sequence 50, Appl
86	42.6	37	10	US-11-055-093-114	Sequence 114, Appl
86	42.6	37	10	US-11-055-093-116	Sequence 116, Appl
86	42.6	37	10	US-11-055-093-119	Sequence 119, Appl
85	42.1	37	10	US-11-055-093-113	Sequence 113, Appl
85	42.1	37	10	US-11-055-093-117	Sequence 117, Appl
84	41.6	37	9	US-10-516-768-24	Sequence 24, Appl
84	41.6	37	9	US-10-760-085-38	Sequence 38, Appl
84	41.6	37	10	US-11-055-093-49	Sequence 49, Appl
84	41.6	37	11	US-11-007-772A-36	Sequence 36, Appl
84	41.6	37	11	US-11-288-495-9	Sequence 9, Appl
82	40.6	15	10	US-11-055-093-216	Sequence 216, Appl
81	40.1	37	9	US-10-516-768-23	Sequence 23, Appl
80.5	39.9	59	10	US-11-055-093-25	Sequence 25, Appl
74	36.6	40	9	US-10-516-768-6	Sequence 6, Appl
74	36.6	125	9	US-10-516-768-8	Sequence 8, Appl
71.5	35.4	56	10	US-11-055-093-28	Sequence 28, Appl
71.5	35.4	56	10	US-11-055-093-30	Sequence 30, Appl
71.5	35.4	56	10	US-11-055-093-32	Sequence 32, Appl
71.5	35.4	56	10	US-11-055-093-36	Sequence 36, Appl
71.5	35.4	57	10	US-11-055-093-34	Sequence 34, Appl
71	35.1	37	9	US-10-516-768-16	Sequence 16, Appl
71	35.1	37	11	US-11-288-495-12	Sequence 12, Appl
71	35.1	125	9	US-10-516-768-18	Sequence 18, Appl
70	34.7	37	9	US-10-516-768-30	Sequence 30, Appl
70	34.7	38	9	US-10-516-768-1	Sequence 1, Appl
70	34.7	38	11	US-11-288-495-10	Sequence 10, Appl
70	34.7	39	9	US-10-516-768-2	Sequence 2, Appl
70	34.7	126	9	US-10-516-768-4	Sequence 4, Appl

95 68 33.7 66 10 US-11-055-093-27 Sequence 27, Appl  
96 68 33.7 66 10 US-11-055-093-29 Sequence 29, Appl  
97 68 33.7 66 10 US-11-055-093-31 Sequence 31, Appl  
98 68 33.7 66 10 US-11-055-093-35 Sequence 35, Appl  
99 68 33.7 67 10 US-11-055-093-33 Sequence 33, Appl  
100 67 33.2 37 9 US-10-516-768-12 Sequence 12, Appl

## ALIGNMENTS

## RESULT 1

US-11-055-093-67  
; Sequence 67, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 67  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-67

Query Match 100.0%; Score 202; DB 10; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.1e-22;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 2

US-11-055-093-69  
; Sequence 69, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 69  
; LENGTH: 37  
; TYPE: PRT

; ORGANISM: Homo sapiens  
US-11-055-093-69

Query Match 99.5%; Score 201; DB 10; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.5e-22;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 3

US-11-055-093-75  
; Sequence 75, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 75  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-75

Query Match 99.5%; Score 201; DB 10; Length 37;  
Best Local Similarity 97.3%; Pred. No. 1.5e-22;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 4

US-11-055-093-92  
; Sequence 92, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 92  
; LENGTH: 37

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-92

Query Match      99.5%; Score 201; DB 10; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.5e-22;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 5
US-11-055-093-74
; Sequence 74, Application US/11055093
; Publication No. US20060094652A1
; GENERAL INFORMATION:
; APPLICANT: LEVY, ODILE ESTHER
; APPLICANT: HANLEY, MICHAEL R.
; APPLICANT: JODKA, CAROLYN M.
; APPLICANT: LEWIS, DIANA Y.
; APPLICANT: SOARES, CHRISTOPHER J.
; APPLICANT: GHOSH, SOUMITRA S.
; APPLICANT: D'SOUZA, LAWRENCE
; APPLICANT: PARKES, DAVID
; APPLICANT: MACK, CHRISTINE M.
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
; FILE REFERENCE: 18528.740
; CURRENT APPLICATION NUMBER: US/11/055,093
; PRIOR FILING DATE: 2005-02-11
; PRIOR FILING DATE: 2004-02-11
; NUMBER OF SEQ ID NOS: 288
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 74
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-74

Query Match      97.5%; Score 197; DB 10; Length 36;
Best Local Similarity 100.0%; Pred. No. 5.4e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 6
US-11-055-093-85
; Sequence 85, Application US/11055093
; Publication No. US20060094652A1
; GENERAL INFORMATION:
; APPLICANT: LEVY, ODILE ESTHER
; APPLICANT: HANLEY, MICHAEL R.
; APPLICANT: JODKA, CAROLYN M.
; APPLICANT: LEWIS, DIANA Y.
; APPLICANT: SOARES, CHRISTOPHER J.
; APPLICANT: GHOSH, SOUMITRA S.
; APPLICANT: D'SOUZA, LAWRENCE
; APPLICANT: PARKES, DAVID
; APPLICANT: MACK, CHRISTINE M.
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
; FILE REFERENCE: 18528.740
; CURRENT APPLICATION NUMBER: US/11/055,093
; PRIOR FILING DATE: 2005-02-11
; PRIOR FILING DATE: 2004-02-11
; NUMBER OF SEQ ID NOS: 288
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 85

; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-85

Query Match      96.5%; Score 195; DB 10; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.1e-21;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 7
US-11-055-093-91
; Sequence 91, Application US/11055093
; Publication No. US20060094652A1
; GENERAL INFORMATION:
; APPLICANT: LEVY, ODILE ESTHER
; APPLICANT: HANLEY, MICHAEL R.
; APPLICANT: JODKA, CAROLYN M.
; APPLICANT: LEWIS, DIANA Y.
; APPLICANT: SOARES, CHRISTOPHER J.
; APPLICANT: GHOSH, SOUMITRA S.
; APPLICANT: D'SOUZA, LAWRENCE
; APPLICANT: PARKES, DAVID
; APPLICANT: MACK, CHRISTINE M.
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
; FILE REFERENCE: 18528.740
; CURRENT APPLICATION NUMBER: US/11/055,093
; PRIOR FILING DATE: 2005-02-11
; PRIOR FILING DATE: 2004-02-11
; NUMBER OF SEQ ID NOS: 288
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 91
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-91

Query Match      96.5%; Score 195; DB 10; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.1e-21;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
   |||||
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 8
US-11-288-495-34
; Sequence 34, Application US/11288495
; Publication No. US20060079448A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Erlandsson, Rikard
; APPLICANT: Frisen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Kristina
; APPLICANT: Hagblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortesman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Osscinak, Amina
; APPLICANT: Patrone, Cesare
; APPLICANT: Ronnholm, Harriet
```



```
/ APPLICANT: LEWIS, DIANA Y.
/ APPLICANT: SOARES, CHRISTOPHER J.
/ APPLICANT: GHOSH, SOUMITRA S.
/ APPLICANT: D'SOUZA, LAWRENCE
/ APPLICANT: PARKES, DAVID
/ APPLICANT: MACK, CHRISTINE M.
/ TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
/ FILE REFERENCE: 18528.740
/ CURRENT APPLICATION NUMBER: US/11/055,093
/ CURRENT FILING DATE: 2005-02-11
/ PRIOR APPLICATION NUMBER: 60/543,407
/ PRIOR FILING DATE: 2004-02-11
/ NUMBER OF SEQ ID NOS: 288
/ SOFTWARE: PatentIn Ver. 3.3
/ SEQ ID NO 73
/ LENGTH: 36
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-11-055-093-73

Query Match          93.6%; Score 189; DB 10; Length 36;
Best Local Similarity 97.2%; Pred. No. 7.9e-21;
Matches 35; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 13
US-11-055-093-84
/ Sequence 84, Application US/11055093
/ Publication No. US20060094652A1
/ GENERAL INFORMATION:
/ APPLICANT: LEVY, ODILE ESTHER
/ APPLICANT: HANLEY, MICHAEL R.
/ APPLICANT: JODKA, CAROLYN M.
/ APPLICANT: LEWIS, DIANA Y.
/ APPLICANT: SOARES, CHRISTOPHER J.
/ APPLICANT: GHOSH, SOUMITRA S.
/ APPLICANT: D'SOUZA, LAWRENCE
/ APPLICANT: PARKES, DAVID
/ APPLICANT: MACK, CHRISTINE M.
/ TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
/ FILE REFERENCE: 18528.740
/ CURRENT APPLICATION NUMBER: US/11/055,093
/ CURRENT FILING DATE: 2005-02-11
/ PRIOR APPLICATION NUMBER: 60/543,407
/ PRIOR FILING DATE: 2004-02-11
/ NUMBER OF SEQ ID NOS: 288
/ SOFTWARE: PatentIn Ver. 3.3
/ SEQ ID NO 84
/ LENGTH: 36
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-11-055-093-84

Query Match          93.6%; Score 189; DB 10; Length 36;
Best Local Similarity 97.2%; Pred. No. 7.9e-21;
Matches 35; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 14
US-11-055-093-89
/ Sequence 89, Application US/11055093
/ Publication No. US20060094652A1
/ GENERAL INFORMATION:
/ APPLICANT: LEVY, ODILE ESTHER
/ APPLICANT: HANLEY, MICHAEL R.
```

```
/ APPLICANT: JODKA, CAROLYN M.
/ APPLICANT: LEWIS, DIANA Y.
/ APPLICANT: SOARES, CHRISTOPHER J.
/ APPLICANT: GHOSH, SOUMITRA S.
/ APPLICANT: D'SOUZA, LAWRENCE
/ APPLICANT: PARKES, DAVID
/ APPLICANT: MACK, CHRISTINE M.
/ TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
/ FILE REFERENCE: 18528.740
/ CURRENT APPLICATION NUMBER: US/11/055,093
/ CURRENT FILING DATE: 2005-02-11
/ PRIOR APPLICATION NUMBER: 60/543,407
/ PRIOR FILING DATE: 2004-02-11
/ NUMBER OF SEQ ID NOS: 288
/ SOFTWARE: PatentIn Ver. 3.3
/ SEQ ID NO 89
/ LENGTH: 37
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-11-055-093-89

Query Match          93.1%; Score 188; DB 10; Length 37;
Best Local Similarity 94.6%; Pred. No. 1.1e-20;
Matches 35; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 15
US-11-055-093-44
/ Sequence 44, Application US/11055093
/ Publication No. US20060094652A1
/ GENERAL INFORMATION:
/ APPLICANT: LEVY, ODILE ESTHER
/ APPLICANT: HANLEY, MICHAEL R.
/ APPLICANT: JODKA, CAROLYN M.
/ APPLICANT: LEWIS, DIANA Y.
/ APPLICANT: SOARES, CHRISTOPHER J.
/ APPLICANT: GHOSH, SOUMITRA S.
/ APPLICANT: D'SOUZA, LAWRENCE
/ APPLICANT: PARKES, DAVID
/ APPLICANT: MACK, CHRISTINE M.
/ TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
/ FILE REFERENCE: 18528.740
/ CURRENT APPLICATION NUMBER: US/11/055,093
/ CURRENT FILING DATE: 2005-02-11
/ PRIOR APPLICATION NUMBER: 60/543,407
/ PRIOR FILING DATE: 2004-02-11
/ NUMBER OF SEQ ID NOS: 288
/ SOFTWARE: PatentIn Ver. 3.3
/ SEQ ID NO 44
/ LENGTH: 37
/ TYPE: PRT
/ ORGANISM: Rattus sp.
US-11-055-093-44

Query Match          92.6%; Score 187; DB 10; Length 37;
Best Local Similarity 91.9%; Pred. No. 1.6e-20;
Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37

RESULT 16
US-11-055-093-86
/ Sequence 86, Application US/11055093
/ Publication No. US20060094652A1
/ GENERAL INFORMATION:
/ APPLICANT: LEVY, ODILE ESTHER
```

; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 86  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-86

Query Match 92.6%; Score 187; DB 10; Length 37;  
Best Local Similarity 91.9%; Pred. No. 1.6e-20;  
Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNGLGVLPTNVGSNTY 37  
|||||

## RESULT 17

US-11-055-093-70  
; Sequence 70, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 70  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-70

Query Match 92.1%; Score 186; DB 10; Length 37;  
Best Local Similarity 94.6%; Pred. No. 2.2e-20;  
Matches 35; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||

## RESULT 18

US-11-055-093-82  
; Sequence 82, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:

; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 82  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-82

Query Match 92.1%; Score 186; DB 10; Length 37;  
Best Local Similarity 94.6%; Pred. No. 2.2e-20;  
Matches 35; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNGLGVLPTNVGSNTY 37  
|||||

## RESULT 19

US-11-055-093-96  
; Sequence 96, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 96  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-96

Query Match 92.1%; Score 186; DB 10; Length 37;  
Best Local Similarity 89.2%; Pred. No. 2.2e-20;  
Matches 33; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNGLGVLPTNVGSNTY 37  
|||||

## RESULT 20

US-11-055-093-189  
; Sequence 189, Application US/11055093  
; Publication No. US20060094652A1



; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 189  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-189

Query Match 92.1%; Score 186; DB 10; Length 37;  
Best Local Similarity 94.6%; Pred. No. 2.2e-20;  
Matches 35; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KNTATCATQRLANFLVHSSNNFGALLPSTNVGSNTY 37  
|||||

RESULT 21  
US-11-055-093-87  
; Sequence 87, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 87  
; LENGTH: 36  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-87

Query Match 90.1%; Score 182; DB 10; Length 36;  
Best Local Similarity 91.7%; Pred. No. 8.3e-20;  
Matches 33; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
QY 2 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KNTATCATQRLANFLVHSSNNLGPVLPSTNVGSNTY 36  
|||||

RESULT 22  
US-11-055-093-71  
; Sequence 71, Application US/11055093

; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 71  
; LENGTH: 36  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-71

Query Match 89.6%; Score 181; DB 10; Length 36;  
Best Local Similarity 94.4%; Pred. No. 1.2e-19;  
Matches 34; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
QY 2 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KNTATCATQRLANFLVRSSNNFGPILPSTNVGSNTY 36  
|||||

RESULT 23  
US-11-055-093-90  
; Sequence 90, Application US/11055093  
; Publication No. US20060094652A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOUMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 90  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-90

Query Match 89.1%; Score 180; DB 10; Length 37;  
Best Local Similarity 91.9%; Pred. No. 1.7e-19;  
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KNTATCATQRLANFLVRSSNNLGPILPSTNVGSNTY 37  
|||||

RESULT 24  
US-11-055-093-88

; Sequence 88, Application US/11055093  
; Publication No. US2006009452A1  
; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 88  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-88

Query Match 88.6%; Score 179; DB 10; Length 37;  
Best Local Similarity 89.2%; Pred. No. 2.3e-19;  
Matches 33; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 25  
US-10-516-768-26  
; Sequence 26, Application US/10516768  
; Publication No. US20050256302A1  
; GENERAL INFORMATION:  
; APPLICANT: MINAMINO, NAOTO  
; APPLICANT: KATAFUCHI, TAKESHI  
; TITLE OF INVENTION: NOVEL PEPTIDES HAVING CAMP PRODUCING ACTIVITY  
; FILE REFERENCE: 62273(71526)  
; CURRENT APPLICATION NUMBER: US/10/516,768  
; CURRENT FILING DATE: 2004-12-03  
; PRIOR APPLICATION NUMBER: PCT/JP03/06641  
; PRIOR FILING DATE: 2003-05-28  
; PRIOR APPLICATION NUMBER: JP 2002-162797  
; PRIOR FILING DATE: 2002-06-04  
; NUMBER OF SEQ ID NOS: 52  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 26  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
; FEATURE:  
; OTHER INFORMATION: C-term amidated  
US-10-516-768-26

Query Match 88.1%; Score 178; DB 9; Length 37;  
Best Local Similarity 91.9%; Pred. No. 3.3e-19;  
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 37

RESULT 26  
US-11-055-093-45  
; Sequence 45, Application US/11055093  
; Publication No. US20060094652A1

; GENERAL INFORMATION:  
; APPLICANT: LEVY, ODILE ESTHER  
; APPLICANT: HANLEY, MICHAEL R.  
; APPLICANT: JODKA, CAROLYN M.  
; APPLICANT: LEWIS, DIANA Y.  
; APPLICANT: SOARES, CHRISTOPHER J.  
; APPLICANT: GHOSH, SOMITRA S.  
; APPLICANT: D'SOUZA, LAWRENCE  
; APPLICANT: PARKES, DAVID  
; APPLICANT: MACK, CHRISTINE M.  
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES  
; FILE REFERENCE: 18528.740  
; CURRENT APPLICATION NUMBER: US/11/055,093  
; CURRENT FILING DATE: 2005-02-11  
; PRIOR APPLICATION NUMBER: 60/543,407  
; PRIOR FILING DATE: 2004-02-11  
; NUMBER OF SEQ ID NOS: 288  
; SOFTWARE: PatentIn Ver. 3.3  
; SEQ ID NO 45  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-11-055-093-45

Query Match 88.1%; Score 178; DB 10; Length 37;  
Best Local Similarity 91.9%; Pred. No. 3.3e-19;  
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 37

## RESULT 27

US-11-033-105A-22  
; Sequence 22, Application US/11033105A  
; Publication No. US20060045853A1  
; GENERAL INFORMATION:  
; APPLICANT: Gebbink, Martijn F.B.G.  
; APPLICANT: Bouma, Barend  
; APPLICANT: Kranenburg, Onno W.  
; APPLICANT: Kroon, Louise M.J.  
; TITLE OF INVENTION: Cross-Beta structure comprising amyloid-binding proteins and  
; TITLE OF INVENTION: methods for detection of the cross-Beta structure, for modulation  
; TITLE OF INVENTION: cross-Beta structures fibril formation and for modulating  
; TITLE OF INVENTION: cross-Beta structure-mediated toxicity  
; FILE REFERENCE: P57716US00  
; CURRENT APPLICATION NUMBER: US/11/033,105A  
; CURRENT FILING DATE: 2005-01-10  
; PRIOR APPLICATION NUMBER: EP 02077797.5  
; PRIOR FILING DATE: 2002-07-09  
; PRIOR APPLICATION NUMBER: PCT/NL2003/000501  
; PRIOR FILING DATE: 2003-07-08  
; NUMBER OF SEQ ID NOS: 45  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 22  
; LENGTH: 37  
; TYPE: PRT  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: Peptide fl-hIAPP from Homo sapiens  
US-11-033-105A-22

Query Match 88.1%; Score 178; DB 11; Length 37;  
Best Local Similarity 91.9%; Pred. No. 3.3e-19;  
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 37

## RESULT 28

```
US-11-288-495-7
; FILE REFERENCE: 21882-517 CIP
; Sequence 7, Application US/11288495
; Publication No. US20060079448A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Erlandsson, Rikard
; APPLICANT: Friesen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Kristina
; APPLICANT: Haggblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortessman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Ossoinak, Amina
; APPLICANT: Patrone, Cesare
; APPLICANT: Ronnholm, Harriet
; APPLICANT: Wikstrom, Lillian
; APPLICANT: Zachrisson, Olof
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
; FILE REFERENCE: 21882-517 CIP
; CURRENT APPLICATION NUMBER: US/11/288,495
; CURRENT FILING DATE: 2005-11-28
; PRIOR APPLICATION NUMBER: US 10/718,071
; PRIOR FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: US 60/427,912
; PRIOR FILING DATE: 2002-11-20
; NUMBER OF SEQ ID NOS: 66
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 7
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-288-495-7

Query Match      88.1%; Score 178; DB 11; Length 37;
Best Local Similarity 91.9%; Pred. No. 3.3e-19;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
    |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGAILSTNVGSNTY 37

RESULT 29
US-11-288-495-35
; Sequence 35, Application US/11288495
; Publication No. US20060079448A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Friesen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Kristina
; APPLICANT: Haggblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortessman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Ossoinak, Amina
; APPLICANT: Patrone, Cesare
; APPLICANT: Ronnholm, Harriet
; APPLICANT: Wikstrom, Lillian
; APPLICANT: Zachrisson, Olof
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
; FILE REFERENCE: 21882-517 CIP
; CURRENT APPLICATION NUMBER: US/11/288,495
; CURRENT FILING DATE: 2005-11-28
; PRIOR APPLICATION NUMBER: US 10/718,071
; PRIOR FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: US 60/427,912
; PRIOR FILING DATE: 2002-11-20
; NUMBER OF SEQ ID NOS: 66
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 7
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-288-495-7

Query Match      88.1%; Score 178; DB 11; Length 37;
Best Local Similarity 91.9%; Pred. No. 3.3e-19;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
    |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGAILSTNVGSNTY 37

RESULT 29
US-11-288-495-35
; Sequence 35, Application US/11288495
; Publication No. US20060079448A1
; GENERAL INFORMATION:
; APPLICANT: Bertilsson, Goran
; APPLICANT: Friesen, Jonas
; APPLICANT: Haegerstrand, Anders
; APPLICANT: Heidrich, Jessica
; APPLICANT: Hellstrom, Kristina
; APPLICANT: Haggblad, Johan
; APPLICANT: Jansson, Katarina
; APPLICANT: Kortessman, Jarkko
; APPLICANT: Lindquist, Per
; APPLICANT: Lundh, Hanna
; APPLICANT: McGuire, Jacqueline
; APPLICANT: Mercer, Alex
; APPLICANT: Nyberg, Karl
; APPLICANT: Ossoinak, Amina
; APPLICANT: Patrone, Cesare
; APPLICANT: Ronnholm, Harriet
; APPLICANT: Wikstrom, Lillian
; APPLICANT: Zachrisson, Olof
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
```

```
FILE REFERENCE: 21882-517 CIP
; CURRENT APPLICATION NUMBER: US/11/288,495
; CURRENT FILING DATE: 2005-11-28
; PRIOR APPLICATION NUMBER: US 10/718,071
; PRIOR FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: US 60/427,912
; PRIOR FILING DATE: 2002-11-20
; NUMBER OF SEQ ID NOS: 66
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 35
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-288-495-35

Query Match      88.1%; Score 178; DB 11; Length 37;
Best Local Similarity 91.9%; Pred. No. 3.3e-19;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
    |||||
Db 1 KCNTATCATQRLANFLVHSSNNFGAILSTNVGSNTY 37

RESULT 30
US-11-145-463-93
; Sequence 93, Application US/11145463
; Publication No. US20060052301A1
; GENERAL INFORMATION:
; APPLICANT: SLEMESH, Ronen
; TITLE OF INVENTION: SPICE VARIANTS OF PEPTIDE YY, NEUROPEPTIDE Y, PANCREATIC PEPTID-
; FILE REFERENCE: 85189-9000
; CURRENT APPLICATION NUMBER: US/11/145,463
; CURRENT FILING DATE: 2005-06-02
; PRIOR APPLICATION NUMBER: US 60/576,414
; PRIOR FILING DATE: 2004-06-03
; PRIOR APPLICATION NUMBER: US 60/672,987
; PRIOR FILING DATE: 2005-04-20
; NUMBER OF SEQ ID NOS: 124
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 93
; LENGTH: 89
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-145-463-93

Query Match      88.1%; Score 178; DB 11; Length 89;
Best Local Similarity 91.9%; Pred. No. 8.8e-19;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
    |||||
Db 34 KCNTATCATQRLANFLVHSSNNFGAILSTNVGSNTY 70

RESULT 31
US-11-030-300-21
; Sequence 21, Application US/11030300
; Publication No. US20060057671A1
; GENERAL INFORMATION:
; APPLICANT: PAN, TAO
; APPLICANT: ORSER, CINDY S.
; TITLE OF INVENTION: IMMOBILIZED PROBES AND METHODS OF DETECTING
; FILE REFERENCE: ADL-102
; CURRENT APPLICATION NUMBER: US/11/030,300
; CURRENT FILING DATE: 2005-01-07
; PRIOR APPLICATION NUMBER: 60/608,541
; PRIOR FILING DATE: 2004-09-10
; NUMBER OF SEQ ID NOS: 31
; SOFTWARE: PatentIn ver. 3.3
; SEQ ID NO 21
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```
; LENGTH: 89
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-030-300-21

Query Match      88.1%; Score 178; DB 11; Length 89;
Best Local Similarity 91.9%; Pred. No. 8.8e-19;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    |||||
Db 34 KNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 70
    |||||

RESULT 32
US-11-145-463-95
; Sequence 95, Application US/11145463
; Publication No. US20060052301A1
; GENERAL INFORMATION:
; APPLICANT: SHEMESH, Ronen
; TITLE OF INVENTION: SPICE VARIANTS OF PEPTIDE YV, NEUROPEPTIDE Y, PANCREATIC PEPTIDE
; FILE REFERENCE: 85189-9000
; CURRENT APPLICATION NUMBER: US/11/145.463
; PRIOR FILING DATE: 2005-06-02
; PRIOR APPLICATION NUMBER: US 60/576,414
; PRIOR FILING DATE: 2004-06-03
; PRIOR APPLICATION NUMBER: US 60/672,987
; PRIOR FILING DATE: 2005-04-20
; NUMBER OF SEQ ID NOS: 124
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 95
; LENGTH: 103
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-145-463-95

Query Match      88.1%; Score 178; DB 11; Length 103;
Best Local Similarity 91.9%; Pred. No. 1e-18;
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    |||||
Db 48 KNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 84
    |||||

RESULT 33
US-11-055-093-68
; Sequence 68, Application US/11055093
; Publication No. US20060094652A1
; GENERAL INFORMATION:
; APPLICANT: LEVY, ODILE ESTHER
; APPLICANT: HANLEY, MICHAEL R.
; APPLICANT: JODKA, CAROLYN M.
; APPLICANT: LEWIS, DIANA Y.
; APPLICANT: SOARES, CHRISTOPHER J.
; APPLICANT: GHOSH, SOUMITRA S.
; APPLICANT: D'SOUZA, LAWRENCE
; APPLICANT: PARKES, DAVID
; APPLICANT: MACK, CHRISTINE M.
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
; FILE REFERENCE: 18528-740
; CURRENT APPLICATION NUMBER: US/11/055,093
; PRIOR FILING DATE: 2005-02-11
; PRIOR APPLICATION NUMBER: US/10/760,085
; PRIOR FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: 60/543,407
; NUMBER OF SEQ ID NOS: 288
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 68
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-68

Query Match      85.6%; Score 173; DB 10; Length 36;
Best Local Similarity 91.7%; Pred. No. 1.7e-18;
Matches 33; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    |||||
Db 1 CNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 36
    |||||

RESULT 34
US-11-055-093-77
; Sequence 77, Application US/11055093
; Publication No. US20060094652A1
; GENERAL INFORMATION:
; APPLICANT: LEVY, ODILE ESTHER
; APPLICANT: HANLEY, MICHAEL R.
; APPLICANT: JODKA, CAROLYN M.
; APPLICANT: LEWIS, DIANA Y.
; APPLICANT: SOARES, CHRISTOPHER J.
; APPLICANT: GHOSH, SOUMITRA S.
; APPLICANT: D'SOUZA, LAWRENCE
; APPLICANT: PARKES, DAVID
; APPLICANT: MACK, CHRISTINE M.
; TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH SELECTABLE PROPERTIES
; FILE REFERENCE: 18528-740
; CURRENT APPLICATION NUMBER: US/11/055,093
; PRIOR FILING DATE: 2005-02-11
; PRIOR APPLICATION NUMBER: 60/543,407
; PRIOR FILING DATE: 2004-02-11
; NUMBER OF SEQ ID NOS: 288
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 77
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-055-093-77

Query Match      85.6%; Score 173; DB 10; Length 36;
Best Local Similarity 91.7%; Pred. No. 1.7e-18;
Matches 33; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    |||||
Db 1 CNTATCATQRLANFLVHSSNFGAILSTNVGSNTY 36
    |||||

RESULT 35
US-10-760-085-8
; Sequence 8, Application US/10760085
; Publication No. US20060051879A9
; GENERAL INFORMATION:
; APPLICANT: Hubert K"ster
; APPLICANT: Daniel Paul Little
; APPLICANT: Suhaib Mahmood Siddiqi
; APPLICANT: Matthew Peter Grealish
; APPLICANT: Subramaniam Marappan
; APPLICANT: Chester Frederick Hassman III
; APPLICANT: Ping Yip
; TITLE OF INVENTION: Capture Compounds, Collections Thereof
; TITLE OF INVENTION: And Methods For Analyzing The Proteome And Complex
; FILE REFERENCE: 24743-2309
; CURRENT APPLICATION NUMBER: US/10/760,085
; PRIOR FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: 60/441,398
; PRIOR FILING DATE: 2003-01-16
; NUMBER OF SEQ ID NOS: 149
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 37
; TYPE: PRT
; ORGANISM: Homo sapien
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GenCore version 5.1.8  
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OM protein - protein search, using sw model

Run on: May 12, 2006, 15:18:26 ; Search time 38 Seconds  
(without alignments)  
93.685 Million cell updates/sec

Title: US-08-870-762B-1

Perfect score: 202

Sequence: 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 100 summaries

Database :

PIR 80:\*

1: Pirl:\*

2: Pirl2:\*

3: Pirl3:\*

4: Pirl4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	187	92.6	93	1	TCRTIA
2	187	92.6	93	1	islet amyloid poly
3	184	91.1	37	2	S05037
4	184	91.1	32	2	insulinoma amyloid
5	178	88.1	89	1	islet amyloid prot
6	171	84.7	89	2	islet amyloid poly
7	169	83.7	89	2	islet amyloid poly
8	160	79.2	91	2	islet amyloid poly
9	153	75.7	135	2	A36118
10	151	74.8	92	2	islet amyloid poly
11	96	47.5	23	2	D33542
12	96	47.5	125	2	I46933
13	93	46.0	29	2	TCCHRP
14	91	45.0	23	2	A61509
15	86	42.6	72	2	I37232
16	86	42.6	127	2	A25864
17	85	42.1	37	2	JH0709
18	85	42.1	128	1	TCRTTR
19	84	41.6	128	1	TCCHUR
20	84	41.6	128	2	B44173
21	84	41.6	134	2	A44173
22	61.5	30.4	166	2	T33970
23	56.5	28.0	430	2	T05980
24	55	27.2	1051	2	T22179
25	55	27.2	140	2	S55259
26	54.5	27.0	140	2	JU0041
27	54	26.7	444	2	A85915
28	54	26.7	444	2	F91070
29	54	26.7	917	2	E96807

30	54	26.7	917	2	S35228	nitrate reductase
31	52.5	26.0	233	2	H72648	hypothetical prote
32	52	25.7	5147	1	IJFPTM	cadherin-related t
33	51.5	25.5	388	2	S57526	cellulase - Fibrob
34	51.5	25.5	436	2	A37953	transcription regu
35	51.5	25.5	638	2	B35816	transcription regu
36	51.5	25.5	649	1	B37953	transcription regu
37	51.5	25.5	649	2	A35816	transcription regu
38	51.5	25.5	1143	2	B84431	probable receptor
39	51	25.2	499	2	S46660	weta protein - Pen
40	51	25.2	555	1	RGASMA	regulatory protein
41	50.5	25.0	424	2	H84806	hypothetical prote
42	50	24.8	110	2	A55991	effector cell prot
43	50	24.8	337	2	A53041	effector cell prot
44	49.5	24.5	458	2	B33560	IL protein - human
45	49	24.3	243	2	B89977	hypothetical prote
46	49	24.3	294	2	T08408	transcription fact
47	49	24.3	359	2	S17905	hypothetical prote
48	49	24.3	1162	2	B97852	hypothetical prote
49	48.5	24.0	405	2	S66525	prostanoid recepto
50	48.5	24.0	844	2	T38730	probable helicase
51	48.5	24.0	1208	2	T27822	hypothetical prote
52	48	23.8	146	1	HGGLB	hemoglobin beta ch
53	48	23.8	146	1	HGGLB	hemoglobin beta ch
54	48	23.8	219	2	JQ1778	SalF3L protein pre
55	48	23.8	256	2	F86856	hypothetical prote
56	48	23.8	599	2	S71134	UV-endonuclease -
57	48	23.8	621	2	S35092	plakoglobin - mous
58	48	23.8	636	2	D82679	hypothetical prote
59	48	23.8	1350	2	T30341	zinc finger protei
60	48	23.8	1687	2	T30244	phosphodiesterase
61	48	23.8	1719	2	T30174	exoribonuclease, v
62	48	23.8	1812	2	I49350	breast/ovarian can
63	48	23.8	2543	2	F69679	polyketide synthas
64	47.5	23.5	69	2	T44123	hypothetical prote
65	47.5	23.5	119	2	D71020	hypothetical prote
66	47.5	23.5	302	2	A32801	fimbrial adhesin p
67	47.5	23.5	320	2	S09208	chorion protein s3
68	47.5	23.5	718	2	A82352	iron(III) compound
69	47	23.3	146	1	HBGSC	hemoglobin beta ch
70	47	23.3	146	1	HBGSC	hemoglobin beta ch
71	47	23.3	146	1	HBGSI	hemoglobin beta ch
72	47	23.3	146	1	HBGSS	hemoglobin beta ch
73	47	23.3	146	1	HBWS	hemoglobin beta ch
74	47	23.3	147	1	HFCHR	hemoglobin rho cha
75	47	23.3	164	2	G75371	hypothetical prote
76	47	23.3	189	2	D97064	spore coat protein
77	47	23.3	309	2	A83702	inosine-uridine nu
78	47	23.3	324	2	F84913	probable FCA-relat
79	47	23.3	324	2	T08729	RING zinc finger p
80	47	23.3	326	1	JQ1238	zinc finger protei
81	47	23.3	366	2	T18820	hypothetical prote
82	47	23.3	388	2	T32307	hypothetical prote
83	47	23.3	470	2	D90323	hypothetical prote
84	47	23.3	470	2	F90347	hypothetical prote
85	47	23.3	517	2	T43358	hnf-3/forhead tra
86	47	23.3	535	2	T32139	hypothetical prote
87	47	23.3	539	2	T38927	hypothetical prote
88	47	23.3	594	2	T05544	hypothetical prote
89	47	23.3	604	2	T31042	hypothetical prote
90	47	23.3	1450	2	T30273	hypothetical prote
91	46.5	23.0	623	2	T40991	probable lysosom
92	46.5	23.0	727	2	T15294	hypothetical prote
93	46.5	23.0	1306	2	S72620	probable reverse t
94	46.5	23.0	1209	2	H86650	protein T3P18.3 [i
95	46.5	23.0	1787	2	AGI360	probable tape-meas
96	46.5	23.0	1788	2	AH1447	probable tape-meas
97	46.5	23.0	1896	2	T08851	Down syndrome cell
98	46	22.8	122	2	T15766	hypothetical prote
99	46	22.8	149	2	T49200	hypothetical prote
100	46	22.8	176	2	AC0207	probable exported

## ALIGNMENTS

## RESULT 1

CJ3542  
N/Alternate names: insulin amyloid precursor - mouse  
N/Alternate names: insulin amyloid precursor  
C/Species: Mus musculus (house mouse)  
C/Date: 10-Sep-1999 #sequence\_revision 10-Sep-1999 #text\_change 09-Jul-2004  
C/Accession: C33542; S05039  
R/Nishi, M.; Chan, S.U.; Nagamatsu, S.; Bell, G.I.; Steiner, D.F.  
Proc. Natl. Acad. Sci. U.S.A. 86, 5738-5742, 1989  
A/Title: Conservation of the sequence of islet amyloid polypeptide in five mammals is conserved  
A/Reference number: A33542; MUID:89345542; PMID:2668946  
A/Accession: C33542  
A/Status: preliminary  
A/Molecule type: mRNA  
A/Residues: 1-93 <NIS>  
A/Cross-references: UNIPROT:P12968; UNIPARC:UPI0000012D0CA; GB:M25389; NID:G194066; PIDN:  
A/Note: the authors translated the codon CTG for residue 18 as Ser, and CTC for residue  
R/Betscholtz, C.; Christman, L.; Engstrom, U.; Rorsman, F.; Svensson, V.; Johnson, K  
FEBS Lett. 251, 261-264, 1989  
A/Title: Sequence divergence in a specific region of islet amyloid polypeptide (IAPP) ex  
A/Reference number: S05037; MUID:89325677; PMID:2666169  
A/Accession: S05039  
A/Status: not compared with conceptual translation  
A/Molecule type: mRNA  
A/Residues: 38-74 <BET>  
A/Cross-references: UNIPARC:UPI000003519C  
A/Note: the authors found both the full length protein and a short form in the pancreas  
C/Superfamily: calcitonin  
C/Keywords: amyloid

Query Match 92.6%; Score 187; DB 1; Length 93;  
Best Local Similarity 91.9%; Pred. No. 1.7e-18;  
Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 38 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 74  
|||||

## RESULT 2

TCRTIA  
N/Alternate names: amylin precursor - rat  
N/Alternate names: amylin precursor; diabetes-associated peptide; insulinoma amyloid pro  
C/Species: Rattus norvegicus (Norway rat)  
C/Date: 31-Dec-1992 #sequence\_revision 31-Dec-1992 #text\_change 09-Jul-2004  
C/Accession: S13566; A30312; B33542; S05038; S05038; B35481  
R/van Mansfeld, A.D.M.; Mosselman, S.; Hoepfener, J.W.M.; Zandberg, J.; van Teeffelen, H  
Biochim. Biophys. Acta 1087, 235-240, 1990  
A/Title: Islet amyloid polypeptide: structure and upstream sequences of the IAPP gene in  
A/Reference number: S13566; MUID:91027936; PMID:2223885  
A/Accession: S13566  
A/Molecule type: DNA  
A/Residues: 1-93 <VAN>  
A/Cross-references: UNIPROT:P12969; UNIPARC:UPI000012D0CA; EMBL:X52820; NID:G56394; PIDN:  
R/Lefvert, J.D.; Newgard, C.B.; Okamoto, H.; Milburn, J.L.; Luskey, K.L.  
Proc. Natl. Acad. Sci. U.S.A. 86, 3127-3130, 1989  
A/Title: Rat amylin: cloning and tissue-specific expression in pancreatic islets.  
A/Reference number: A30312; MUID:89240689; PMID:2654937  
A/Accession: A30312  
A/Molecule type: mRNA  
A/Residues: 1-93 <LEP>  
A/Cross-references: UNIPARC:UPI000012D0CA; GB:J04544; NID:G202887; PIDN:AAA40730.1; PID:  
R/Nishi, M.; Chan, S.U.; Nagamatsu, S.; Bell, G.I.; Steiner, D.F.  
Proc. Natl. Acad. Sci. U.S.A. 86, 5738-5742, 1989  
A/Title: Conservation of the sequence of islet amyloid polypeptide in five mammals is co  
A/Reference number: A33542; MUID:89345542; PMID:2668946  
A/Accession: B33542  
A/Molecule type: mRNA  
A/Residues: 1-93 <NIS>  
A/Cross-references: UNIPARC:UPI000012D0CA; GB:M25390; NID:G204676; PIDN:AAA41359.1; PID:  
R/Asai, J.; Nakazato, M.; Kangawa, K.; Matsukura, S.; Matsuo, H.

Biochem. Biophys. Res. Commun. 164, 400-405, 1989  
A/Title: Isolation and sequence determination of rat islet amyloid polypeptide.  
A/Reference number: A33426; MUID:90026410; PMID:2679555  
A/Accession: A33426  
A/Molecule type: protein  
A/Residues: 38-74 <ASA>

A/Cross-references: UNIPARC:UPI000003519C  
R/Betscholtz, C.; Christman, L.; Engstrom, U.; Rorsman, F.; Svensson, V.; Johnson, K  
FEBS Lett. 251, 261-264, 1989  
A/Title: Sequence divergence in a specific region of islet amyloid polypeptide (IAPP) ex  
A/Reference number: S05037; MUID:89325677; PMID:2666169  
A/Accession: S05038  
A/Status: not compared with conceptual translation  
A/Molecule type: mRNA  
A/Residues: 38-74 <BET>  
A/Cross-references: UNIPARC:UPI000003519C  
R/Asai, J.; Nakazato, M.; Miyazato, K.; Matsuo, H.; Matsukura, S.  
Biochem. Biophys. Res. Commun. 169, 788-795, 1990  
A/Title: Regional distribution and molecular forms of rat islet amyloid polypeptide.  
A/Reference number: A35481; MUID:90290528; PMID:2357234  
A/Accession: A35481  
A/Molecule type: protein  
A/Residues: 38-74 <AS2>  
A/Cross-references: UNIPARC:UPI000003519C  
A/Accession: B35481  
A/Molecule type: protein  
A/Residues: 56-74 <AS3>  
A/Cross-references: UNIPARC:UPI000003519C  
A/Note: the authors found both the full length protein and a short form in the pancreas  
C/Superfamily: calcitonin  
C/Keywords: amyloid; signal sequence #status predicted <SIG>  
E/1-23/Domain: signal sequence #status predicted <SIG>  
E/24-37/Domain: amino-terminal propeptide #status predicted <PRO1>  
E/38-74/Domain: insulinoma amyloid protein #status experimental <MAT>  
F/76-93/Domain: carboxyl-terminal propeptide #status predicted <PRO2>  
F/39-44/Disulfide bonds: #status predicted  
F/74/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 92.6%; Score 187; DB 1; Length 93;  
Best Local Similarity 91.9%; Pred. No. 1.7e-18;  
Matches 34; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 38 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 74  
|||||

## RESULT 3

S05037  
insulinoma amyloid protein - Chinese hamster  
C/Species: Cricetulus griseus (Chinese hamster)  
C/Date: 07-Sep-1990 #sequence\_revision 07-Sep-1990 #text\_change 09-Jul-2004  
C/Accession: S05037  
R/Betscholtz, C.; Christman, L.; Engstrom, U.; Rorsman, F.; Svensson, V.; Johnson, K  
FEBS Lett. 251, 261-264, 1989  
A/Title: Sequence divergence in a specific region of islet amyloid polypeptide (IAPP) ex  
A/Reference number: S05037; MUID:89325677; PMID:2666169  
A/Accession: S05037  
A/Status: not compared with conceptual translation  
A/Molecule type: mRNA  
A/Residues: 1-37 <BET>  
A/Cross-references: UNIPROT:P19890; UNIPARC:UPI00000466B2  
C/Superfamily: calcitonin  
C/Keywords: amyloid

Query Match 91.1%; Score 184; DB 2; Length 37;  
Best Local Similarity 89.2%; Pred. No. 1.7e-18;  
Matches 33; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||

## RESULT 4

S1316  
islet amyloid protein precursor - golden hamster  
C:Species: Mesocricetus auratus (golden hamster)  
C:Date: 21-Nov-1993 #sequence\_revision 10-Nov-1995 #text\_change 09-Jul-2004  
C:Accession: S13116  
R:Nishi, M.; Bell, G.I.; Steiner, D.F.  
Nucleic Acids Res. 18, 6726, 1990  
A:Title: Sequence of a cDNA encoding Syrian hamster islet amyloid polypeptide precursor.  
A:Reference number: S13116; MUID:91067499; PMID:2251153  
A:Accession: S13116  
A:Status: preliminary  
A:Molecule type: mRNA  
A:Residues: 1-92 <NIS>  
A:Cross-references: UNIPROT:P23442; UNIPARC:UPI000012D0C4; EMBL:X56067; NID:g49633; PIDN  
C:Superfamily: calcitonin

Query Match 91.1%; Score 184; DB 2; Length 92;  
Best Local Similarity 89.2%; Pred. No. 4.4e-18;  
Matches 33; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNPGLPTPTNVGSNTY 37

|||||  
Db 37 KCNTATCATQRLANFLVHSSNNLGPVLSPTNVGSNTY 73

## RESULT 5

TCHUIA  
islet amyloid polypeptide precursor [validated] - human  
N:Alternate names: amylin precursor; diabetes-associated peptide; insulinoma amyloid pro  
C:Species: Homo sapiens (man)  
C:Date: 31-Mar-1993 #sequence\_revision 31-Mar-1993 #text\_change 09-Jul-2004  
A:Accession: S04016; JCI1408; A41406; B32116; B32116; A34499; A26619; A39985; A60061; B60  
R:Mosselman, S.; Hoepfner, J.W.M.; Lips, C.J.M.; Jansz, H.S.  
FEBS Lett. 247, 154-158, 1989  
A:Title: The complete islet amyloid polypeptide precursor is encoded by two exons.  
A:Reference number: S04016; MUID:89211434; PMID:2651160  
A:Accession: S04016  
A:Molecule type: DNA  
A:Residues: 1-89 <MOS>  
A:Cross-references: UNIPROT:P10957; UNIPARC:UPI000012D0C4; EMBL:X14902  
R:Hoepfner, J.W.M.; Oosterwijk, C.; Visser-Vernooij, H.J.; Lips, C.J.M.; Jansz, H.S.  
Biochem. Biophys. Res. Commun. 189, 1569-1577, 1992  
A:Title: Characterization of the human islet amyloid polypeptide/amylin gene transcripts  
A:Reference number: JCI1408; MUID:93129228; PMID:1282806  
A:Accession: JCI1408  
A:Molecule type: DNA  
A:Residues: 1-89 <HOE>  
A:Cross-references: UNIPARC:UPI000012D0C4; EMBL:X68830; NID:g32582; PIDN:CAA48724.1; PID  
R:Nishi, M.; Sanke, T.; Saino, S.; Eddy, R.L.; Fan, Y.S.; Byers, M.G.; Shows, T.B.; Bell  
Mol. Endocrinol. 3, 1775-1781, 1989  
A:Title: Human islet amyloid polypeptide gene: complete nucleotide sequence, chromosomal  
A:Reference number: A41406; MUID:90114181; PMID:2608057  
A:Accession: A41406  
A:Molecule type: DNA  
A:Residues: 1-89 <NIS>  
A:Cross-references: UNIPARC:UPI000012D0C4; GB:M26650; NID:g184047; PIDN:AAA35983.1; PID:  
R:Sanke, T.; Bell, G.I.; Sample, C.; Rubenstein, A.H.; Steiner, D.F.  
J. Biol. Chem. 263, 17243-17246, 1988  
A:Title: An islet amyloid peptide is derived from an 89-amino acid precursor by proteoly  
A:Reference number: A92691; MUID:89034238; PMID:3053705  
A:Accession: A32116  
A:Molecule type: mRNA  
A:Residues: 1-89 <SAN>  
A:Cross-references: UNIPARC:UPI000012D0C4; GB:J04422; NID:g178621; PIDN:AAAS2281.1; PID:  
A:Accession: B32116  
A:Molecule type: mRNA  
A:Residues: 27-89 <SA2>  
A:Cross-references: UNIPARC:UPI000016AAMD6  
A>Note: the authors obtained five independent clones for this mRNA variant, which may re  
R:Roberts, A.N.; Leighton, B.; Todd, J.A.; Cockburn, D.; Schofield, F.N.; Sutton, R.; Ho  
Proc. Natl. Acad. Sci. U.S.A. 86, 9662-9666, 1989

A:Title: Molecular and functional characterization of amylin, a peptide associated with  
A:Reference number: A34499; MUID:90099324; PMID:2690069  
A:Accession: A34499  
A:Molecule type: DNA  
A:Residues: 30-89 <ROB>  
A:Cross-references: UNIPARC:UPI0000173497  
R:Westermarck, P.; Wernstedt, C.; Wilander, E.; Hayden, D.W.; O'Brien, T.D.; Johnson, K.I  
Proc. Natl. Acad. Sci. U.S.A. 84, 3881-3885, 1987  
A:Title: Amyloid fibrils in human insulinoma and islets of Langerhans of the diabetic c  
A:Reference number: A26619; MUID:87231921; PMID:3035556  
A:Accession: A26619  
A:Molecule type: protein  
A:Residues: 34-68, 'X', '70 <WES>  
A:Cross-references: UNIPARC:UPI000002C9BF  
R:Cooper, G.J.S.; Willis, A.C.; Clark, A.; Turner, R.C.; Sim, R.B.; Reid, K.B.M.  
Proc. Natl. Acad. Sci. U.S.A. 84, 8628-8632, 1987  
A:Title: Purification and characterization of a peptide from amyloid-rich pancreases of  
A:Reference number: A39985; MUID:88068642; PMID:3317417  
A:Accession: A39985  
A:Molecule type: protein  
A:Residues: 34-70 <COO>  
A:Cross-references: UNIPARC:UPI000002B886  
R:Nakazato, M.; Asai, J.; Miyazato, M.; Matsukura, S.; Kangawa, K.; Matsuo, H.  
Regul. Pept. 31, 179-186, 1990  
A:Title: Isolation and identification of islet amyloid polypeptide in normal human panc  
A:Reference number: A60061; MUID:91219694; PMID:2091067  
A:Accession: A60061  
A:Molecule type: protein  
A:Residues: 34-70 <NAK>  
A:Cross-references: UNIPARC:UPI000002B886  
A:Experimental source: normal pancreas  
A:Accession: B60061  
A:Molecule type: protein  
A:Residues: 50-70 <NA2>  
A:Cross-references: UNIPARC:UPI0000173498  
A:Experimental source: normal pancreas  
R:Besholtz, C.; Svensson, V.; Rorsman, F.; Engstroem, U.; Westermark, G.T.; Wilander, P  
Exp. Cell Res. 183, 484-493, 1989  
A:Title: Islet amyloid polypeptide (IAPP): cDNA cloning and identification of an amylo  
A:Reference number: A60599; MUID:89356900; PMID:2670595  
A:Accession: A60599  
A:Status: nucleic acid sequence not shown; not compared with conceptual translation  
A:Molecule type: mRNA  
A:Residues: 1-89 <BET>  
A:Cross-references: UNIPARC:UPI000012D0C4  
R:Christman, L.; Rorsman, F.; Stenman, G.; Westermark, P.; Betsholtz, C.  
FEBS Lett. 267, 160-166, 1990  
A:Title: The human islet amyloid polypeptide (IAPP) gene. Organization, chromosomal loca  
A:Reference number: S10729; MUID:90306394; PMID:2365085  
A:Accession: S10729  
A:Status: preliminary  
A:Molecule type: DNA  
A:Residues: 1-89 <CHR>  
A:Cross-references: UNIPARC:UPI000012D0C4  
R:van Mansfeld, A.D.M.; Mosselman, S.; Hoepfner, J.W.M.; Zandberg, J.; van Teeffelen, H  
Biochim. Biophys. Acta 1087, 235-240, 1990  
A:Title: Islet amyloid polypeptide: structure and upstream sequences of the IAPP gene in  
A:Reference number: S13566; MUID:91027936; PMID:2223885  
A:Accession: S13567  
A:Status: preliminary; translation not shown  
A:Molecule type: DNA  
A:Residues: 1-89 <VAN>  
A:Cross-references: UNIPARC:UPI000012D0C4; EMBL:X52818; NID:g32589; PIDN:CAA37002.1; PID  
R:Mosselman, S.; Hoepfner, J.W.M.; Zandberg, J.; van Mansfeld, A.D.M.; Geurts van Kesse  
FEBS Lett. 239, 227-232, 1988  
A:Title: Islet amyloid polypeptide: identification and chromosomal localization of the I  
A:Reference number: S01779; MUID:89031237; PMID:3181427  
A:Accession: S01779  
A:Molecule type: DNA  
A:Residues: 28-89 <MO2>  
A:Cross-references: UNIPARC:UPI000016A552; EMBL:X13859; NID:g32584; PIDN:CAB57803.1; PID  
R:Westermarck, P.; Wernstedt, C.; Wilander, E.; Sletten, K.  
Biochem. Biophys. Res. Commun. 140, 827-831, 1986

A;Title: A novel peptide in the calcitonin gene related peptide family as an amyloid fibril precursor  
A;Reference number: A36385; MUID:87048863; PMID:3535798  
A;Accession: A36385  
A;Molecule type: protein  
A;Residues: 34; 'X', 36-39, 'S', 41-52 <WE2>  
A;Cross-references: UNIPARC:UPI0000173499  
R;Cort, J.; Liu, Z.; Lee, G.; Harris, S.M.; Prickett, K.S.; Gaeta, L.S.L.; Andersen, N.H. Biochem. Biophys. Res. Commun. 204, 1088-1095, 1994  
A;Title: beta-Structure in human amylin and two designer beta-peptides: CD and NMR spectra  
A;Reference number: PC2383; MUID:95071438; PMID:7980582  
A;Contents: annotation; circular dichroism and NMR studies  
C;Comments: This protein is a major component of the islet amyloid deposited in the pancreas as a hormone.  
C;Genetics:  
A;Gene: GDB:IAPP  
A;Cross-references: GDB:120074; OMIM:147940  
A;Map position: 12p12.3-12p12.1  
A;Introns: 27/2  
C;Superfamily: calcitonin  
C;Keywords: amidated carboxyl end; amyloid; pancreatic islet; type 2 diabetes  
F;1-22/Domain: signal sequence #status predicted <SIG>  
F;23-33/Domain: amino-terminal propeptide #status predicted <PRO1>  
F;34-70/Product: islet amyloid polypeptide #status experimental <MAT>  
F;50-70/Product: islet amyloid polypeptide (17-37) #status experimental <MAT2>  
F;53-62/Domain: amyloid fibril-forming #status experimental <FIB>  
F;72-89/Domain: carboxyl-terminal propeptide #status predicted <PRO2>  
F;35-40/Disulfide bonds: #status predicted  
F;70/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 88.1%; Score 178; DB 1; Length 89;  
Best Local Similarity 91.9%; Pred. No. 2.8e-17;  
Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPIPLPTNVGSNTY 37  
|||||  
Db 34 KCNTATCATQRLANFLVHSSNNFGAILSTNVGSNTY 70  
|||||

RESULT 6  
A33542  
islet amyloid polypeptide precursor - cat  
C;Species: Felis silvestris catus (domestic cat)  
C;Date: 21-Feb-1990 #sequence revision 21-Feb-1990 #text\_change 09-Jul-2004  
A;Accession: A33542; A60499; E26619  
R;Nishi, M.; Chan, S.J.; Nagamatsu, S.; Bell, G.I.; Steiner, D.F. Proc. Natl. Acad. Sci. U.S.A. 86, 5738-5742, 1989  
A;Title: Conservation of the sequence of islet amyloid polypeptide in five mammals is compared  
A;Reference number: A33542; MUID:89345542; PMID:2668946  
A;Accession: A33542  
A;Molecule type: mRNA  
A;Residues: 1-89 <NIS>  
A;Cross-references: UNIPROT:P12967; UNIPARC:UPI000012D0C3; GB:M25388; NID:g163861; PIDN:R;Besholtz, C.; Christmanson, L.; Engstroem, U.; Rorsman, F.; Jordan, K.; O'Brien, T.D. Diabetes 39, 118-122, 1990  
A;Title: Structure of cat islet amyloid polypeptide and identification of amino acid residues  
A;Reference number: A60499; MUID:91006862; PMID:2210054  
A;Accession: A60499  
A;Status: not compared with conceptual translation  
A;Molecule type: mRNA  
A;Residues: 34-70 <BET>  
A;Cross-references: UNIPARC:UPI000003519A  
R;Westermarck, P.; Wernstedt, C.; Willander, E.; Hayden, D.W.; O'Brien, T.D.; Johnson, K.H. Proc. Natl. Acad. Sci. U.S.A. 84, 3881-3885, 1987  
A;Title: Amyloid fibrils in human insulinoma and islets of Langerhans of the diabetic cat  
A;Reference number: A36619; MUID:87231921; PMID:3035556  
A;Accession: B26619  
A;Status: preliminary  
A;Molecule type: protein  
A;Residues: 34; 'X', 36-39, 'X', 41-60 <WES>  
A;Cross-references: UNIPARC:UPI00000269C0  
C;Superfamily: calcitonin  
C;Keywords: amidated carboxyl end; amyloid; pancreas

F;70/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 84.7%; Score 171; DB 2; Length 89;  
Best Local Similarity 86.5%; Pred. No. 2.5e-16;  
Matches 32; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPIPLPTNVGSNTY 37  
|||||  
Db 34 KCNTATCATQRLANFLVHSSNNLGAAILSTNVGSNTY 70  
|||||

RESULT 7  
S22344  
islet amyloid polypeptide precursor - dog  
N;Alternate names: amylin precursor; IAPP; insulinoma amyloid protein  
C;Species: Canis lupus familiaris (dog)  
C;Date: 22-Nov-1993 #sequence\_revision 01-Dec-1995 #text\_change 09-Jul-2004  
C;Accession: S22344; A35476  
R;Albrandt, K.; Mull, E.; Cooper, G.J.S.; Johnson, M.J. Biochim. Biophys. Acta 1130, 97-99, 1992  
A;Title: Nucleotide sequence of a cDNA for canine amylin.  
A;Reference number: S22344; MUID:92182022; PMID:1543754  
A;Accession: S22344  
A;Molecule type: mRNA  
A;Residues: 1-89 <ALB>  
A;Cross-references: UNIPROT:P17716; UNIPARC:UPI000012D0C1; EMBL:X59998; NID:g870; PIDN:C;Jordan, K.; Murrcaugh, M.P.; O'Brien, T.D.; Westermarck, P.; Besholtz, C.; Johnson, K.H. Biochem. Biophys. Res. Commun. 169, 502-508, 1990  
A;Title: Canine IAPP cDNA sequence provides important clues regarding diabetogenesis and  
A;Reference number: A35476; MUID:90290487; PMID:2192709  
A;Accession: A35476  
A;Molecule type: mRNA  
A;Residues: 43-66; 'T', 68 <JOR>  
A;Cross-references: UNIPARC:UPI000016C3DA; GB:M37720; NID:g163960; PIDN:AAA30849.1; PID:C;Superfamily: calcitonin

Query Match 83.7%; Score 169; DB 2; Length 89;  
Best Local Similarity 86.5%; Pred. No. 4.8e-16;  
Matches 32; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPIPLPTNVGSNTY 37  
|||||  
Db 34 KCNTATCATQRLANFLVHSSNNLGAAILSTNVGSNTY 70  
|||||

RESULT 8  
A36118  
islet amyloid polypeptide precursor - degu  
C;Species: Octodon degus (degu)  
C;Date: 28-Mar-1991 #sequence\_revision 28-Mar-1991 #text\_change 09-Jul-2004  
C;Accession: A36118  
R;Nishi, M.; Steiner, D.F. Mol. Endocrinol. 4, 1192-1198, 1990  
A;Title: Cloning of complementary DNAs encoding islet amyloid polypeptide, insulin, and  
A;Reference number: A36118; MUID:91155953; PMID:2293024  
A;Accession: A36118  
A;Status: preliminary  
A;Molecule type: mRNA  
A;Residues: 1-91 <NIS>  
A;Cross-references: UNIPROT:P22889; UNIPARC:UPI000012D0C7; GB:M57669; NID:g202469; PIDN:C;Superfamily: calcitonin

Query Match 79.2%; Score 160; DB 2; Length 91;  
Best Local Similarity 81.1%; Pred. No. 8.4e-15;  
Matches 30; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPIPLPTNVGSNTY 37  
|||||  
Db 37 KCNTATCATQRLNFLVHSSNNLGAAILPTNVGSNTY 73  
|||||

RESULT 9  
A56855

islet amyloid polypeptide precursor - chicken  
N;Alternate names: IAPP  
C;Species: Gallus gallus (chicken)  
C;Date: 11-Aug-1995 #sequence\_revision 11-Aug-1995 #text\_change 09-Jul-2004  
C;Accession: A56855  
R;Fan, L.; Westermarck, G.; Chan, S. J.; Steiner, D. F.  
Mol. Endocrinol. 8, 713-721, 1994  
A;Title: Altered gene structure and tissue expression of islet amyloid polypeptide in the  
A;Reference number: A56855; MUID:95021303; PMID:7935487  
A;Accession: A56855  
A;Status: preliminary  
A;Molecule type: mRNA; DNA  
A;Residues: 1-135 <FAN>  
A;Cross-references: UNIPROT.Q90743; UNIPARC.UPI00000FDB07; GB:L16955; NID:g289789; PIDN:  
C;Superfamily: calcitonin  
C;Keywords: hormone

Query Match 75.7%; Score 153; DB 2; Length 135;  
Best Local Similarity 78.4%; Pred. No. 1.1e-13;  
Matches 29; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

Qy	1 KCNTATCATQRLANFLVHSSNNFGLPPTNVGSNTY 37       
Db	80 KCNTATCVTORLADFLVRSSNIGAIYSPTNVGSNTY 116       :             :

RESULT 10  
D33542  
islet amyloid polypeptide precursor - guinea pig  
C:Species: *Cavia porcellus* (guinea pig)  
C:date: 21-Feb-1990 #sequence\_revision 04-Sep-1992 #text\_change 09-Jul-2004  
C:Accession: D33542

R; Nishi, M.; Chan, S.J.; Nagamatsu, S.; Bell, G.I.; Steiner, D.F. Proc Natl. Acad. Sci. U.S.A. 86, 5738-5742, 1989

A; Title: Conservation of the sequence of islet amyloid polypeptide in five mammals is compared

A; Reference number: A33542; MUID:89345542; PMID:2668946

A; Accession: D33542

A; Status: preliminary

A; Molecule type: mRNA

A; Residues: 1-92 <NTS>

A; Cross-references: UNIPROT:P12966; UNIPARC:UPI000012D0C2; GB:M25387; NID:g191271; PIDN:

C; Note: the authors translated the codon CTA for residue 87 as Cys

C; Superfamily: calcitonin

```

Query Match          74.8%; Score 151; DB 2; Length 92;
Best Local Similarity 78.4%; Pred. No. 1.4e-13;
Matches 29; Conservative 2; Mismatches 6; Indels 0; Gaps 0;

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Qy	37
<div style="background-color: black; height: 1em;"></div>	
1 KCNTATCATQRLANFLVHSSNNFGLPPTNVGSNTY	
<div style="background-color: black; height: 1em;"></div>	
:	
<div style="background-color: black; height: 1em;"></div>	
Db	73
<div style="background-color: black; height: 1em;"></div>	
37 KCNTATCATQRLTNFLVRSSHNLGAALLPTDVGSNYY	
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RESULT 11  
I46933  
islet amyloid polypeptide - European hare (fragment)  
C:Species: Lepus capensis europaeus (European hare)  
C:Date: 21-Feb-1997 #sequence\_revision 21-Feb-1997 #text\_change 09-Jul-2004  
C:Accession: I46933  
R:Christmansson, L.; Betscholtz, C.; Leckstrom, A.; Engstrom, U.; Cortie, C.; Johnson, K.H.  
D:Diabetologia 36, 183-189, 1993  
A:Title: Islet amyloid polypeptide in the rabbit and European hare: studies on its relation to diabetes mellitus  
A:Reference number: I46933; MUID:93215963; PMID:8462765  
A:Accession: I46933  
A>Status: preliminary; translated from GB/EMBL/DBJ  
A:Molecule type: mRNA  
A:Residues: 1-23 <CHR>  
A:Cross-references: UNIPROT:Q07333; UNIPARC:UPI000012D0C5; GB:S57802; NID:g299056; PIDN:C  
C:Superfamily: calcitonin

Query Match 47.5%; Score 96; DB 2; Length 23;  
Best Local Similarity 78.3%; Pred. NO. 1.2e-06;  
Matches 18; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

Qy 9 TQRLANFLVHSSNFGPILPPTN 31  
| | | | | : | | | | |  
Db 1 TQRLANFLIHSSNFGAIFSPN 23

RESULT 12  
TCCHRP  
calcitonin gene-related peptide precursor - chicken  
C/Species: Gallus gallus (Chicken)  
C/Date: 30-Jun-1991 #sequence revision 30-Jun-1991 #text\_change 09-Jul-2004  
C/Accession: S00154, I50183; A24855  
R/Mintzelle, S.; Cressent, M.; Delchaye, M.C.; Second, N.; Milhaud, G.; Jul  
FEBS Lett. 223, 63-68, 1987  
A/Title: Sequence and expression of the chicken calcitonin gene.  
A/Reference number: S00153; MUID:89030046; PMID:3656142

A;Accession: S00134  
A;Molecule type: DNA  
A;Residues: 1-125 <MIN>  
A;Cross-references: UNIPROT:P10286; UNIPARC:UPI0000126E2F; EMBL:X063111  
A;Note: the sequences of codons 31-33 and 34-38 are interchanged in the  
A;Note: the authors translated the codon GAC for residue 56 as Glu  
R;asmolseqs, F.; Jullienne, A.; Day, F.; Minvielle, S.; Milhaud, G.; Moreau  
EMBO J. 4, 2603-2607, 1985  
A;Title: Elucidation of the nucleotide sequence of chicken calcitonin  
A;Reference number: A25725; MUID:86030240; PMID:4054101

A:Contents: annotation  
R:Minvielle, S.; Cressent, M.; Lasmoles, F.; Jullienne, A.; Milhaud, G.; Moukhtar, M.S. FEBS Lett. 203, 7-10, 1986  
A:Title: Isolation and partial characterization of the calcitonin gene in a lower vertebrate  
A:Reference number: I50183; PMID:86248126; PMID:3487468  
A:Accession: I50183  
A:Status: preliminary; translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 74-125 <MF2>  
A:Cross-references: UNIPARC:UPI00001712BC; GB:D00007; NID:G222801; PIDN:BAA00006.1; PID:G222801  
C:Comment: The calcitonin gene codes for two mRNA species by tissue-specific alternative splicing. The calcitonin gene-related peptide (CGRP) is a 27-residue peptide that acts as a peripheral nervous system codes for calcitonin gene-related peptide.

A;introns: 29/2; 73/2  
 C;Superfamily: calcitonin  
 C;Keywords: alternative splicing; amidated carboxyl end; neuropeptide  
 F;80-116/Product: calcitonin gene-related peptide #status predicted <MAT>  
 F;81-86/Disulfide bonds: #status predicted  
 F;116/Modified site: amidated carboxyl end (Phe) (amide in mature form from fol

Qy 2 CNTATCATQRLANFLVHS-----SNNFGPIIPPTNVGSNTY 37  
||||| : ||| : ||| : ||| :  
Db 81 CNTATCTWHLADFLSRSGVGKNPF----VPTNVGSKAF 116

```

RESULT 13
A61509
islet amyloid polypeptide - cougar (fragment)
C/Species: Felis concolor (cougar)
C/Date: 19-Mar-1997 #sequence_revision 26-Feb-1998 #text_change 11-May-2000
C/Accession: A61509
R/Johnson, K.H.; Wernstedt, C.; O'Brien, T.D.; Westermark, P.
Comp. Biochem. Physiol. B 98, 115-119, 1991
A/Title: Amyloid in the pancreatic islets of the cougar (Felis concolor) is
A/Reference number: A61509; MUID:91284578; PMID:2060275
A/Accession: A61509
A/Molecule type: protein
A/Residues: 1-29 <JOH>
A/Cross-references: UNIPARC:UP1000017660F
C/Superfamily: calcitonin

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Query Match 46.0%; Score 93; DB 2; Length 29;  
Best Local Similarity 74.1%; Pred. No. 3.8e-06;

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Matches 20; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

Qy 1 KNTATCATQRLANFLVHSSNFGPIL 27
    ||||| ||||| ||||| ||||| |||||
Db 1 KXNTATXATQRLANFLIRSSXNLGAIL 27

RESULT 14
I46934
A:Title: amyloid polypeptide - rabbit (fragment)
C:Species: Oryctolagus cuniculus (domestic rabbit)
C>Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 09-Jul-2004
C:Accession: I46934
R:Christmannson, L.; Betsholtz, C.; Leckstrom, A.; Engstrom, U.; Cortie, C.; Johnson, K.H.
Diabetologia 36, 183-188, 1993
A:Title: Islet amyloid polypeptide in the rabbit and European hare: studies on its relat
A:Reference number: I46933; MUID:93215963; PMID:8462765
A:Accession: I46934
A>Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: mRNA
A:Residues: 1-23 <CHR>
A:Cross-references: UNIPROT:Q07334; UNIPARC:UPI000016C61B; GB:S57804; NID:g299058; PIDN:
C:Superfamily: calcitonin

Query Match 45.0%; Score 91; DB 2; Length 23;
Best Local Similarity 81.0%; Pred. No. 5.7e-06;
Matches 17; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

Qy 9 TORLANFLVHSSNFGPILPP 29
    ||||| ||||| ||||| ||||| |||||
Db 1 TORLANFLVHSSNFGALFSP 21

RESULT 15
I37232
A:Title: calcitonin gene-related peptide 2 - human (fragment)
C:Species: Homo sapiens (man)
C>Date: 06-Sep-1996 #sequence_revision 06-Sep-1996 #text_change 09-Jul-2004
C:Accession: I37232
R:Steenbergh, P.H.; Hoppener, J.W.; Zandberg, J.; Lips, C.J.; Jansz, H.S.
FEBS Lett. 183, 403-407, 1985
A:Title: A second human calcitonin/CGRP gene.
A:Reference number: I37232; MUID:85180007; PMID:2985435
A:Accession: I37232
A>Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: mRNA
A:Residues: 1-72 <RES>
A:Cross-references: UNIPROT:P10092; UNIPARC:UPI000016A6C1; EMBL:X02404; NID:g29933; PIDN:
C:Superfamily: calcitonin

Query Match 42.6%; Score 86; DB 2; Length 72;
Best Local Similarity 50.0%; Pred. No. 8.9e-05;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    ||||| ||||| ||||| ||||| |||||
Db 28 CNTATCVTHRLAGLLSRSGGVMKSNFVPTNVGSKAF 63

RESULT 16
A25864
A:Title: calcitonin gene-related peptide beta precursor - human
N:Alternate names: calcitonin gene-related peptide II
C:Species: Homo sapiens (man)
C>Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C:Accession: A25864; JH0620; B26142; A34565
R:Steenbergh, P.H.; Hoppener, J.W.M.; Zandberg, J.; Visser, A.; Lips, C.J.M.; Jansz, H.
FEBS Lett. 209, 97-103, 1986
A:Title: Structure and expression of the human calcitonin/CGRP genes.
A:Reference number: A25864; MUID:87105923; PMID:3492393
A:Accession: A25864
A:Molecule type: DNA
A:Residues: 1-127 <STE>
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A:Cross-references: UNIPROT:P10092; UNIPARC:UPI0000126E36
R:Kitamura, K.; Kangawa, K.; Kawamoto, M.; Ichiki, Y.; Matsuo, H.; Eto, T.
Biochem. Biophys. Res. Commun. 185, 134-141, 1992
A:Title: Isolation and characterization of peptides which act on rat platelets, from a p
A:Reference number: JH0618; MUID:92287083; PMID:1318039
A:Accession: JH0620
A:Molecule type: protein
A:Residues: 82,'X','84-87,'X','89-104 <KIT>
A:Cross-references: UNIPARC:UPI0000176610
A:Experimental source: pHeochromocytoma
R:Petermann, J.B.; Born, W.; Chang, J.Y.; Fischer, J.A.
J. Biol. Chem. 262, 542-545, 1987
A:Title: Identification in the human central nervous system, pituitary, and thyroid of a
A:Reference number: A92637; MUID:87109142; PMID:3492492
A:Accession: B26142
A:Molecule type: protein
A:Residues: 82,'X','84-87,'X','89-91,'X','93-98,'X','100-105,'X','107-109 <PET>
A:Cross-references: UNIPARC:UPI0000176611
R:Wimalawansa, S.J.; Morris, H.R.; Etienne, A.; Blench, I.; Panico, M.; MacIntyre, I.
Biochem. Biophys. Res. Commun. 167, 993-1000, 1990
A:Title: Isolation, purification and characterization of beta-hCGRP from human spinal co
A:Reference number: A34565; MUID:90211348; PMID:2322288
A:Accession: A34565
A:Molecule type: protein
A:Residues: 82-86;104-117 <WIM>
A:Cross-references: UNIPARC:UPI0000176612; UNIPARC:UPI0000176613
C:Comment: Calcitonin gene-related-peptide II peptide is a potent vasorelaxant.
C:Genetics:
A:Gene: GDB:CALCB; CALC2
A:Cross-references: GDB:120572; OMIM:114160
A:Map position: 11p15.2-11p15.1
C:Superfamily: calcitonin
C:Keywords: amidated carboxyl end; neuropeptide
F:82-118/Product: calcitonin gene-related peptide beta #status experimental <MAT>
F:83-88/Dialfide bonds: #status experimental
F:118/Modified site: amidated carboxyl end (Phe) (amide in mature form from following gl

Query Match 42.6%; Score 86; DB 2; Length 127;
Best Local Similarity 50.0%; Pred. No. 0.00016;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    ||||| ||||| ||||| ||||| |||||
Db 83 CNTATCVTHRLAGLLSRSGGVMKSNFVPTNVGSKAF 118

RESULT 17
JH0709
A:Title: calcitonin gene-related peptide - sheep
C:Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C>Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 09-Jul-2004
C:Accession: JH0709
R:Miyata, A.; Jiang, L.; Minamino, N.; Arimura, A.
Biochem. Biophys. Res. Commun. 187, 1474-1479, 1992
A:Title: Identification of calcitonin gene related peptide in ovine hypothalamic extract
A:Reference number: JH0709; MUID:93038624; PMID:1417824
A:Accession: JH0709
A:Molecule type: protein
A:Residues: 1-37 <MIY>
A:Cross-references: UNIPROT:P30881; UNIPARC:UPI0000035153
A:Experimental source: hypothalamus
C:Comment: This protein has adenylate cyclase stimulating activity.
C:Superfamily: calcitonin
C:Keywords: amidated carboxyl end; neuropeptide
F:37/Modified site: amidated carboxyl end (Phe) #status experimental

Query Match 42.1%; Score 85; DB 2; Length 37;
Best Local Similarity 50.0%; Pred. No. 6.4e-05;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
    ||||| ||||| ||||| ||||| |||||
Db 2 CNTATCVTHRLAGLLSRSGGVMKSNFVPTNVGSKAF 37
```

## RESULT 18

## TCRTR

calcitonin gene-related peptide precursor - rat  
C:Species: Rattus norvegicus (Norway rat)  
C:Date: 19-Feb-1984 #sequence\_revision 19-Feb-1984 #text\_change 09-Jul-2004  
C:Accession: A01524; B22949  
R:Amara, S.G.; Jonas, V.; Rosenfeld, M.G.; Ong, E.S.; Evans, R.M.  
Nature 298, 240-244, 1982  
A:Title: Alternative RNA processing in calcitonin gene expression generates mRNAs encoding a novel peptide  
A:Reference number: A01524; MUID:82220111; PMID:6283379  
A:Accession: A01524  
A:Molecule type: mRNA  
A:Residues: 1-128 <AMA>  
A:Cross-references: UNIPROT:P01256; UNIPARC:UPI00001708AA; GB:L29188; GB:J00714; GB:N0001708AA  
R:Jonas, V.; Lin, C.R.; Kawashima, E.; Semon, D.; Swanson, L.W.; Mermod, J.J.; Evans, R.  
Proc. Natl. Acad. Sci. U.S.A. 82, 1994-1998, 1985  
A:Title: Alternative RNA processing events in human calcitonin/calcitonin gene-related peptide  
A:Reference number: A94030; MUID:85166259; PMID:3872459  
A:Accession: B22949  
A:Molecule type: mRNA  
A:Residues: 1-39, 'A', 40-49, 'L', 50-67, 70-128 <JON>  
A:Cross-references: UNIPARC:UPI0000126E33  
C:Superfamily: calcitonin  
C:Keywords: alternative splicing; amidated carboxyl end; neuropeptide  
F:1-25/Domain: signal sequence #status predicted <SIG>  
F:83-119/Product: calcitonin gene-related peptide #status predicted <MAT>  
F:84-89/Diulfide bonds: #status predicted  
F:119/Modified site: amidated carboxyl end (Phe) (amide in mature form from following gl

Query Match 42.1%; Score 85; DB 1; Length 128;  
Best Local Similarity 50.0%; Pred. No. 0.00022;  
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;  
QY 2 CNTATCATORLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 84 CNTATCTVTRLAAGLSRGGVVKDNFVPTNVGSEAF 119

## RESULT 19

## TCCHUR

calcitonin gene-related peptide alpha precursor [validated] - human  
N:Alternate names: calcitonin gene-related peptide I; CGRP-I  
C:Species: Homo sapiens (man)  
C:Date: 30-Sep-1987 #sequence\_revision 02-Jul-1996 #text\_change 09-Jul-2004  
C:Accession: S07644; A22949; B22716; I55536; J00005; S10813; A26142; JH0619; I52204; I84  
R:Broad, P.M.; Symes, A.J.; Thakker, R.V.; Craig, R.K.  
Nucleic Acids Res. 17, 6999-7011, 1989  
A:Title: Structure and methylation of the human calcitonin/alpha-CGRP gene.  
A:Reference number: S07643; MUID:89386053; PMID:2571128  
A:Accession: S07644  
A:Molecule type: DNA  
A:Residues: 1-128 <BRO>  
A:Cross-references: UNIPROT:P06881; UNIPARC:UPI0000126E30; EMBL:X15943; NID:g29613; PIDN  
A:Note: the authors translated the codon CAG for residue 19 as Glu  
R:Jonas, V.; Lin, C.R.; Kawashima, E.; Semon, D.; Swanson, L.W.; Mermod, J.J.; Evans, R.  
Proc. Natl. Acad. Sci. U.S.A. 82, 1994-1998, 1985  
A:Title: Alternative RNA processing events in human calcitonin/calcitonin gene-related peptide  
A:Reference number: A94030; MUID:85166259; PMID:3872459  
A:Accession: A22949  
A:Molecule type: mRNA  
A:Residues: 1-128 <JON>  
A:Cross-references: UNIPARC:UPI0000126E30; GB:M12667; NID:g179825; PIDN:AAAS1914.1; PID  
R:Edbrooke, M.R.; Parker, D.; McVey, J.H.; Riley, J.H.; Sorenson, G.D.; Pettengill, O.S.  
EMBO J. 4, 715-724, 1985  
A:Title: Expression of the human calcitonin/CGRP gene in lung and thyroid carcinoma.  
A:Reference number: A91034; MUID:85230541; PMID:2408883  
A:Accession: B22716  
A:Molecule type: mRNA  
A:Residues: 'V', 50-75, 'S', 76-128 <EDB>  
A:Cross-references: UNIPARC:UPI000017349A  
R:Steenbergh, P.H.; Hoppener, J.W.; Zandberg, J.; Van de Ven, W.J.; Jansz, H.S.; Lips, O

J. Clin. Endocrinol. Metab. 59, 358-360, 1984  
A:Title: Calcitonin gene related peptide coding sequence is conserved in the human genome  
A:Reference number: I55536; MUID:84240176; PMID:6610687  
A:Accession: I55536  
A>Status: translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 77-128 <RES>  
A:Cross-references: UNIPARC:UPI000016A651; GB:M28637; NID:g180467; PIDN:AAAS2012.1; PID  
R:Morris, H.R.; Panik, M.; Etienne, T.; Tippins, J.; Girgis, S.I.; MacIntyre, I.  
Nature 308, 746-748, 1984  
A:Title: Isolation and characterization of human calcitonin gene-related peptide.  
A:Reference number: A93329; MUID:84191466; PMID:6609312  
A:Accession: J00005  
A:Molecule type: protein  
A:Residues: 83-119 <MOR>  
A:Cross-references: UNIPARC:UPI000002B78E  
A:Note: this peptide was detected in medullary thyroid carcinoma tissue and in plasma  
R:Zaidi, M.; Brain, S.D.; Tippins, J.R.; di Marzo, V.; Moonga, B.S.; Chambers, T.J.; Mo  
Biochem. J. 269, 775-780, 1990  
A:Title: Structure-activity relationship of human calcitonin-gene-related peptide.  
A:Reference number: S10813; MUID:90358780; PMID:2390067  
A:Accession: S10813  
A:Molecule type: protein  
A:Residues: 83-99, 'A', 101-119 <ZAI>  
A:Cross-references: UNIPARC:UPI0000017349B  
R:Petermann, J.B.; Born, W.; Chang, J.Y.; Fischer, J.A.  
J. Biol. Chem. 262, 542-545, 1987  
A:Title: Identification in the human central nervous system, pituitary, and thyroid of a  
A:Reference number: A92637; MUID:87109142; PMID:3492492  
A:Accession: A26142  
A:Molecule type: protein  
A:Residues: 83-88, 'X', 90-101, 'X', 103-111, 'X', 113-115, 'X', 117 <PET>  
A:Cross-references: UNIPARC:UPI000017349C  
R:Kitamura, K.; Kangawa, K.; Kawamoto, M.; Ichiki, Y.; Matsuo, H.; Eto, T.  
Biochem. Biophys. Res. Commun. 185, 134-141, 1992  
A:Title: Isolation and characterization of peptides which act on rat platelets, from a  
A:Reference number: JH0618; MUID:92287083; PMID:1318039  
A:Accession: JH0619  
A:Molecule type: protein  
A:Residues: 83, 'X', 85-88, 'X', 90-108 <KIT>  
A:Cross-references: UNIPARC:UPI0000017349D  
A:Experimental source: pheochromocytoma  
R:Nelkin, B.D.; Rosenfeld, K.I.; de Bustros, A.; Leong, S.S.; Roos, B.A.; Baylin, S.B.  
Biochem. Biophys. Res. Commun. 123, 648-655, 1984  
A:Title: Structure and expression of a gene encoding human calcitonin and calcitonin gen  
A:Reference number: I52204; MUID:85022523; PMID:6148938  
A:Accession: I52204  
A>Status: translated from GB/EMBL/DBJ  
A:Molecule type: mRNA  
A:Residues: 48-119 <RE2>  
A:Cross-references: UNIPARC:UPI000016A6C0; GB:K03512; NID:g180465; PIDN:AAAS2011.1; PID  
R:Craig, R.K.; Riley, J.H.; Edbrooke, M.R.; Broad, P.M.; Foord, S.M.; Al-Kazwini, S.J.;  
Biochem. Soc. Symp. 52, 91-105, 1986  
A:Title: Expression and function of the human calcitonin/alpha-CGRP gene in health and d  
A:Reference number: I39387; MUID:87213363; PMID:3034287  
A:Accession: I84508  
A>Status: translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 77-128 <RE3>  
A:Cross-references: UNIPARC:UPI000016A651; GB:M26094; NID:g179798; PIDN:AAAS1912.1; PID  
C:Comment: This peptide is a potent vasorelaxant.  
C:Comment: This peptide increases the rate and force of contraction of rat auricles in v  
C:Genetics:  
A:Gene: GDB:CALCA; CALCI  
A:Cross-references: GDB:120571; OMIM:114130  
A:Map position: lip15.2-lip15.1  
A:Introns: 29/2; 76/2  
C:Superfamily: calcitonin  
C:Keywords: alternative splicing; amidated carboxyl end; neuropeptide; vasodilator  
F:83-119/Product: calcitonin gene-related peptide alpha #status experimental <CTN>  
F:84-89/Diulfide bonds: #status experimental  
F:119/Modified site: amidated carboxyl end (Phe) (amide in mature form from following gl

```
Query Match      41.6%; Score 84; DB 1; Length 128;
Best Local Similarity 50.0%; Pred. No. 0.0003;
Matches 20; Conservative 2; Mismatches 10; Indels 8; Gaps 2;

Qy 2 CNTATCATORLANFLVHS-----SNNFGPILPPTNVGSNTY 37
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| :
Db 84 CDTATCVTHRLAGLLRSRGGVVKNF-----VPTNVGSKAF 119

RESULT 20
B44173
calcitonin gene-related peptide alpha precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 30-Jan-1993 #sequence_revision 12-Mar-1993 #text_change 09-Jul-2004
C:Accession: B44173
R:Amara, S.G.; Arriza, J.L.; Leff, S.E.; Swanson, L.W.; Evans, R.M.; Rosenfeld, M.G.
Science 229, 1094-1097, 1985
A:Title: Expression in brain of a messenger RNA encoding a novel neuropeptide homologous
A:Reference number: A44173; MUID:85300490; PMID:2994212
A:Accession: B44173
A:Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-128 <ANA>
A:Cross-references: UNIPROT:P01256; UNIPARC:UPI00001708AB; GB:M11597; NID:g203226; PIDN:
A:Note: the authors translated the codon CGC for residue 99 as Ser
C:Superfamily: calcitonin
C:Keywords: neuropeptide

Query Match      41.6%; Score 84; DB 2; Length 128;
Best Local Similarity 50.0%; Pred. No. 0.0003;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

Qy 2 CNTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37
   ||||| ||||| ||||| ||||| ||||| ||||| :
Db 84 CNTATCVTHRLAGLLRSRGGVVKNFVPTNVGSEAF 119

RESULT 21
A44173
calcitonin gene-related peptide beta precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 30-Jan-1993 #sequence_revision 12-Mar-1993 #text_change 09-Jul-2004
C:Accession: A44173; S18300
R:Amara, S.G.; Arriza, J.L.; Leff, S.E.; Swanson, L.W.; Evans, R.M.; Rosenfeld, M.G.
Science 229, 1094-1097, 1985
A:Title: Expression in brain of a messenger RNA encoding a novel neuropeptide homologous
A:Reference number: A44173; MUID:85300490; PMID:2994212
A:Accession: A44173
A:Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-134 <ANA>
A:Cross-references: UNIPROT:P10093; UNIPARC:UPI0000126E39; GB:M11596; NID:g203232; PIDN:
A:Note: the authors translated the codon AAA for residue 47 as Phe, and CGC for residue
R:Wang, M.W.; Young, A.A.; Rink, T.J.; Cooper, G.J.S.
FEBS Lett. 291, 195-198, 1991
A:Title: (8-37)h-CGRP antagonizes actions of amylin on carbohydrate metabolism in vitro
A:Reference number: S18300; MUID:92038032; PMID:1936264
A:Accession: S18300
A:Molecule type: protein
A:Residues: 96-104, 'S', '106-112, 'N', '114-125 <WAN>
A:Cross-references: UNIPARC:UPI000003515C
C:Superfamily: calcitonin
C:Keywords: neuropeptide

Query Match      41.6%; Score 84; DB 2; Length 134;
Best Local Similarity 50.0%; Pred. No. 0.00032;
Matches 18; Conservative 1; Mismatches 17; Indels 0; Gaps 0;

Qy 2 CNTATCATORLANFLVHSSNFGPILPPTNVGSNTY 37
   ||||| ||||| ||||| ||||| ||||| ||||| :
Db 90 CNTATCVTHRLAGLLRSRGGVVKNFVPTNVGSKAF 125
```

## RESULT 22

T33970  
hypothetical protein F46E10.11 - Caenorhabditis elegans  
C:Species: Caenorhabditis elegans  
C>Date: 29-Oct-1999 #sequence\_revision 29-Oct-1999 #text\_change 29-Oct-1999  
C:Accession: T33970  
R:Johnson, D.; Bradshaw, H.  
submitted to the EMBL Data Library, February 1999  
A:Description: The sequence of C. elegans cosmid F46E10.  
A:Reference number: Z21446  
A:Accession: T33970  
A:Status: preliminary; translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 1-166 <JOH>  
A:Cross-references: UNIPARC:UPI000017B9D5; EMBL:AF125955; PIDN:AAD14711.1; GSPDB:GN000023  
A:Experimental source: strain Bristol N2; clone F46E10  
C:Genetics:  
A:Gene: CESP:F46E10.11  
A:Map position: 5  
A:Introns: 55/1

Query Match 30.4%; Score 61.5; DB 2; Length 166;  
Best Local Similarity 36.4%; Pred. No. 0.48;  
Matches 16; Conservative 3; Mismatches 12; Indels 13; Gaps 2;

Qy 2 CNTATCATORLANFLVHSSN-----NFGPILPPTNVGS 34  
 ||||| ||||| ||||| ||||| ||||| ||||| :  
Db 65 CNTATCSTSSYCCQI--RSNWRGCGCRVFNVEPTLPPTTTRS 106

## RESULT 23

T05980  
hypothetical protein F17M5.50 - Arabidopsis thaliana  
C:Species: Arabidopsis thaliana (mouse-ear cress)  
C>Date: 30-Apr-1999 #sequence\_revision 30-Apr-1999 #text\_change 31-Dec-2004  
C:Accession: T05980  
R:Bevan, M.; Rose, M.; Hempel, S.; Entian, K.D.; Hoheisel, J.; Mewes, H.W.; Mayer, K.F.X.  
submitted to the Protein Sequence Database, March 1999  
A:Reference number: Z15263  
A:Accession: T05980  
A:Molecule type: DNA  
A:Residues: 1-430 <BEV>  
A:Cross-references: UNIPROT:Q9SZA6; UNIPARC:UPI00000A24DD; EMBL:AL035678; GSPDB:GN000062;  
A:Experimental source: cultivar Columbia; BAC clone F17M5  
C:Genetics:  
A:Gene: ATSP:F17M5.50  
A:Map position: 4  
C:Superfamily: hypothetical protein containing F-box domain

Query Match 28.0%; Score 56.5; DB 2; Length 430;  
Best Local Similarity 36.4%; Pred. No. 6.2;  
Matches 16; Conservative 5; Mismatches 16; Indels 7; Gaps 2;

Qy 1 KCNTATCATORLANFLVH-----SSNFGPILP-PTNVGSNTY 37  
 ||||| ||||| ||||| ||||| ||||| ||||| :  
Db 211 KGNTWCARENSDGNHIIICFDTRERFGPLLPVNVIDNEY 254

## RESULT 24

T22179  
hypothetical protein F44F1.4 - Caenorhabditis elegans  
C:Species: Caenorhabditis elegans  
C>Date: 15-Oct-1999 #sequence\_revision 15-Oct-1999 #text\_change 09-Jul-2004  
C:Accession: T22179  
R:Dobson, R.  
submitted to the EMBL Data Library, October 1996  
A:Reference number: Z19527  
A:Accession: T22179  
A:Status: preliminary; translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 1-559 <WIL>  
A:Cross-references: UNIPROT:O02263; UNIPARC:UPI000007C6F9; EMBL:Z81083; PIDN:CAB03104.1;  
A:Experimental source: clone F44F1



R.;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; Mayhew, M.W.; Miller, L.; Grobeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; Apodaca, N.; Nature 409, 529-533, 2001

A>Title: Genome sequence of enterohemorrhagic Escherichia coli O157:H7.

A|Reference number: A85480; MUID:21074935; PMID:11206551

A|Accession: A85915

A|Status: Preliminary

A|Molecule type: DNA

A|Residues: 1-444 <STO>

A|Cross-references: UNIPROT:Q8X922; UNIPARC:UPI00001658DC; GB:AE005174; NID:gl2517110; i

A|Experimental source: strain O157:H7, substrain EDL933

C|Genetics:

A|Gene: Z3973

C|Superfamily: hypothetical protein b1439

Query Match 26.7%; Score 54; DB 2; Length 444;  
Best Local Similarity 36.1%; Pred.No.14;  
Matches 13; Conservative 8; Mismatches 15; Indels 0; Gaps 0;

Qy 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNT 36  
Db 395 KANAAGLAVALSDWRIRSEDDGLLLSFTNIVSES 430

RESULT 28

F91070

probable transcription regulator [imported] - Escherichia coli (strain O157:H7, substra

C|Species: Escherichia coli

C|Date: 18-Jul-2001 #sequence\_revision 18-Jul-2001 #text\_change 09-Jul-2004

C|Accession: F91070

C|Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, C.G.  
Sasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.  
DNA Res. 8, 11-22, 2001

A>Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and gen

A|Reference number: A99629; MUID:21156231; PMID:11258796

A|Accession: F91070

A|Status: preliminary

A|Molecule type: DNA

A|Residues: 1-444 <HAV>

A|Cross-references: UNIPROT:Q8X922; UNIPARC:UPI0000D088C; GB:BA0000007; PIDN:BAB36957.1,

A|Experimental source: strain O157:H7, substrain RIMD 0509952

C|Genetics:

A|Gene: EC93534

C|Superfamily: hypothetical protein b1439

Query Match 26.7%; Score 54; DB 2; Length 444;  
Best Local Similarity 36.1%; Pred.No.14;  
Matches 13; Conservative 8; Mismatches 15; Indels 0; Gaps 0;

Qy 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNT 36  
Db 395 KANAAGLAVALSDWRIRSEDDGLLLSFTNIVSES 430

RESULT 29

E96807

nitrate reductase 1 (NR1), 46724-43362 [imported] - Arabidopsis thaliana

C|Species: Arabidopsis thaliana (mouse-ear cress)

C|Date: 02-Mar-2001 #sequence\_revision 02-Mar-2001 #text\_change 09-Jul-2004

C|Accession: E96807

R|Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso,

Chen, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.,  
anzen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 816-820, 2000

A|Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C.  
C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali,  
Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

A|Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon,  
ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A>Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.

A|Reference number: A86141; MUID:21016719; PMID:11130712

A|Accession: E96807

A|Status: preliminary

```
A:Molecule type: DNA
A:Residues: 1-917 <STO>
A:Cross-references: UNIPROT:P11832; UNIPARC:UPI000000011AFA; GB:AB005173; NID:G6437524; PID:
C:Genetics:
A:Gene: T3288.9
A:Map position: 1
C:Superfamily: nitrate reductase (NADH); cytochrome b5 core homology; cytochrome-b5 redu
C:Keywords: heme; iron; metalloprotein
F:580,603/Binding site: heme iron (His) (axial ligands) #status predicted

Query Match      26.7%; Score 54; DB 2; Length 917;
Best Local Similarity 56.2%; Pred. No. 30;
Matches 9; Conservative 3; Mismatches 4; Indels 0; Gaps 0;

Qy      14 NFLVHSSNNFGPIPLPP 29
      | | | | | | | | | |
Db      629 NVSVHGASNFGPLLAP 644

RESULT 30
335228
nitrate reductase (NADH) (EC 1.7.1.1) 1 - Arabidopsis thaliana
C:Species: Arabidopsis thaliana (mouse-ear cress)
C:Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C:Accession: S35228; S01640; S16495; S32018
R:Wilkinson, J.O.; Crawford, N.M.
Mol. Gen. Genet. 239, 289-297, 1993
A:Title: Identification and characterization of a chlorate-resistant mutant of Arabidops
A:Reference number: S35228; MUID:93287999; PMID:8510658
A:Accession: S35228
A:Molecule type: DNA
A:Residues: 1-917 <WTL>
A:Cross-references: UNIPROT:P11832; UNIPARC:UPI0000016DB93; EMBL:Z19050; NID:G22756; PIDN
R:Cheng, C.; Dewdney, J.; Nam, H.; den Boer, B.G.W.; Goodman, H.M.
EMBO J. 7, 3309-3314, 1988
A:Title: A new locus (NIA1) in Arabidopsis thaliana encoding nitrate reductase.
A:Reference number: S01640; MUID:89091069; PMID:2905260
A:Accession: S01640
A:Molecule type: mRNA
A:Residues: 525-917 <CHE1>
A:Cross-references: UNIPARC:UPI000016DB95; EMBL:X13434; NID:G16402; PIDN:CAA31786.1; PID
A:Note: translation of nucleotide sequence not complete
A:Accession: S16495
A>Status: translation not shown
A:Molecule type: DNA
A:Residues: 342-360 <CHE2>
A:Cross-references: UNIPARC:UPI000016DB94; EMBL:X13436; NID:G16401; PIDN:CAA31788.1; PID
C:Genetics:
A:Gene: NIA1
A:Map position: 1
A:Introns: 352/1; 399/1; 476/3
C:Complex: homodimer
C:Superfamily: nitrate reductase (NADH); cytochrome b5 core homology; cytochrome-b5 redu
C:Keywords: chromoprotein; electron transfer; FAD; flavoprotein; heme; homodimer; iron;
F:93-485/Domain: molybdopterin-binding domain homology <PCO>
F:545-619/Domain: cytochrome b5 core homology <CB5>
F:667-917/Domain: cytochrome-b5 reductase homology <CBR>
F:197/Binding site: molybdopterin (Cys) (covalent) #status predicted
F:436/Disulfide bonds: interchain #status predicted
F:580,603/Binding site: heme iron (His) (axial ligands) #status predicted
F:731,889/Binding site: NAD (Lys, Cys) #status predicted
F:771/Binding site: FAD (Tyr) #status predicted

Query Match      26.7%; Score 54; DB 2; Length 917;
Best Local Similarity 56.2%; Pred. No. 30;
Matches 9; Conservative 3; Mismatches 4; Indels 0; Gaps 0;

Qy      14 NFLVHSSNNFGPIPLPP 29
      | | | | | | | | | |
Db      629 NVSVHGASNFGPLLAP 644

RESULT 31
```

```
H72648
hypothetical protein APO622 - Aeropyrum pernix (strain K1)
C:Species: Aeropyrum pernix
C:Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C:Accession: H72648
R:Kawarabayashi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takah
awa, H.; Takaniya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; K
DNA Res. 6, 83-101, 1999
A:Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyr
A:Reference number: A72450; MUID:99310339; PMID:10382966
A:Accession: H72648
A>Status: preliminary
A:Molecule type: DNA
A:Residues: 1-233 <KAW>
A:Cross-references: UNIPROT:Q9YEF4; UNIPARC:UPI000005DC05; DDBJ:AP0000060; NID:G5104188;
A:Experimental source: strain K1
C:Genetics:
A:Gene: APO622

Query Match      26.0%; Score 52.5; DB 2; Length 233;
Best Local Similarity 46.2%; Pred. No. 12;
Matches 12; Conservative 5; Mismatches 8; Indels 1; Gaps 1;

Qy      10 QRLANFLVHS-SNNFGPIPLPTNVGS 34
      : | | | | | | | | | |
Db      28 ERFAAFTTPSVSSNVTMLPPTSIAS 53

RESULT 32
IUFFTM
cadherin-related tumor suppressor precursor - fruit fly (Drosophila melanogaster)
C:Species: Drosophila melanogaster
C:Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Feb-1997
C:Accession: A41087; B41087
R:Mahoney, P.A.; Weber, U.; Onofrechuk, P.; Biesmann, H.; Bryant, P.J.; Goodman, C.S.
Cell 67, 853-868, 1991
A:Title: The fat tumor suppressor gene in Drosophila encodes a novel member of the cadhe
A:Reference number: A41087; MUID:92069752; PMID:1959133
A:Accession: A41087
A:Molecule type: mRNA
A:Residues: 143-485; 1279-5147 <MAH>
A:Cross-references: UNIPARC:UPI000017434B; UNIPARC:UPI000017434C; GB:M80537
A:Accession: B41087
A:Molecule type: DNA
A:Residues: 1-142; 487-1278 <MA2>
A:Cross-references: UNIPARC:UPI000017434D; UNIPARC:UPI000017434E; GB:M80537
A:Note: 1229-Gly and 1233-Ser were also found
C:Genetics:
A:Gene: fat
A:Cross-references: FlyBase:FBgn0001075
C:Superfamily: cadherin-related tumor suppressor; cadherin repeat homology; EGF homology
C:Keywords: calcium binding; cell adhesion; duplication; transmembrane protein
F:1-35/Domain: signal sequence #status predicted <SIG>
F:36-5147/Product: cadherin-related tumor suppressor #status predicted <MAT>
F:36-4583/Domain: extracellular #status predicted <EXT>
F:51-156/Domain: cadherin repeat homology <CR1>
F:159-270/Domain: cadherin repeat homology <CR2>
F:271-382/Domain: cadherin repeat homology <CR3>
F:390-494/Domain: cadherin repeat homology <CR4>
F:497-599/Domain: cadherin repeat homology <CR5>
F:602-708/Domain: cadherin repeat homology <CR6>
F:718-822/Domain: cadherin repeat homology <CR7>
F:831-942/Domain: cadherin repeat homology <CR8>
F:948-1049/Domain: cadherin repeat homology <CR9>
F:1052-1153/Domain: cadherin repeat homology <CR10>
F:1156-1278/Domain: cadherin repeat homology <CR11>
F:1281-1384/Domain: cadherin repeat homology <CR12>
F:1387-1489/Domain: cadherin repeat homology <CR13>
F:1492-1601/Domain: cadherin repeat homology <CR14>
F:1607-1713/Domain: cadherin repeat homology <CR15>
F:1717-1823/Domain: cadherin repeat homology <CR16>
F:1826-1922/Domain: cadherin repeat homology <CR17>
F:1925-2027/Domain: cadherin repeat homology <CR18>
```

RESULT 34  
A37953  
transcription regulator PAN-1 - golden hamster (fragment)  
C/Species: Mesocricetus auratus (golden hamster)  
C/Date: 06-Dec-1991 #sequence\_revision 06-Dec-1991 #text\_change 09-Jul-2004  
C/Accession: A37953  
R/German, M.S.; Bianar, M.A.; Nelson, C.; Moss, L.G.; Rutter, W.J.  
Mol. Endocrinol. 5, 292-299, 1991  
A/Title: Two related helix-loop-helix proteins participate  
A/Reference number: A37953; MUID:91246228; PMID:1710033  
A/Accession: A37953  
A/Status: preliminary; not compared with conceptual translation  
A/Molecule type: mRNA  
A/Residues: 1-436 <GER>  
A/Cross-references: UNIPROT:P98180; UNIPARC:UPI0000176415

```

Query Match      25.5%; Score 51.5; DB 1; Length 649;
Best Local Similarity 70.6%; Pred. No. 46;
Matches 12; Conservative 0; Mismatches 4; Indels 1; Gaps 1;

QY 18 HSSNNFGPILPPTNVGS 34
      ||||| - - - |||
Db 340 HSSNNFSP-SPSPTVGS 355

```

[illegible]

RESULT 45

B89977  
hypochemical protein SA1707 [imported] - Staphylococcus aureus (strain N315)  
C:Species: Staphylococcus aureus  
C:Date: 10-May-2001 #sequence\_revision 10-May-2001 #text\_change 09-Jul-2004  
C:Accession: B89977  
R:Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; Oguni, C.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, K.; C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.  
Lancet 357, 1225-1240, 2001  
A:Title: Whole genome sequencing of methicillin-resistant Staphylococcus aureus.  
A:Reference number: A89759; MUID:21311952; PMID:11418146  
A:Accession: B89977  
A>Status: preliminary  
A:Molecule type: DNA  
A:Residues: 1-243 <KUR>  
A:Cross-references: UNIPROT:Q99S25; UNIPARC:UPI00000D77A4; GB:BA000018; PID:gl3701684; 1  
A:Experimental source: strain N315  
C:Genetics:  
A:Gene: SA1707

Query Match	24.3%;	Score 49;	DB 2;	Length 243;
Best Local Similarity	41.4%;	Pred. NO. 37;		
Matches	12;	Conservative	3;	Mismatches 14; Indels 0; Gaps 0;

QY 9 TQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 126 TNRLTGDVIESDTFTGTVGFENHGGRTY 154

RESULT 46

T08408  
transcription factor homolog F18B3.150 - Arabidopsis thaliana  
C:Species: Arabidopsis thaliana (mouse-ear cress)  
C:Date: 11-Jun-1999 #sequence\_revision 11-Jun-1999 #text\_change 09-Jul-2004  
C:Accession: T08408  
R:Quetier, F.; Rieger, M.; Gabel, C.; Mueller-Auer, S.; Schaefer, M.; Zipp, M.; Salanoubat submitted to the Protein Sequence Database, May 1999  
A:Reference number: Z16409  
A:Accession: T08408  
A:Molecule type: DNA  
A:Residues: 1-294 <QUE>  
A:Cross-references: UNIPROT:Q9SVL2; UNIPARC:UPI00000A322F; EMBL:AL049862; GSPDB:GN000061, 1  
A:Experimental source: cultivar Columbia; BAC clone F18B3  
C:Genetics:  
A:Gene: ATSP:F18B3.150  
A:Map position: 3  
A:Introns: 172/3

Query Match	24.3%;	Score 49;	DB 2;	Length 294;
Best Local Similarity	35.3%;	Pred. NO. 44;		
Matches	12;	Conservative	8;	Mismatches 12; Indels 2; Gaps 1;

QY 3 NTATCATQRLANF--LVHSSNNFGPILPPTNVGS 34  
DB 89 STSSGASSCISNFWDLHTKNNNSKTAPYNNVPS 122

RESULT 47

S17905  
hypochemical protein M1 - La France disease virus  
C:Species: La France disease virus  
C:Date: 19-Mar-1997 #sequence\_revision 24-Jul-1997 #text\_change 09-Jul-2004  
C:Accession: S17905  
R:Harmsen, M.C.; Tolner, B.; Kram, A.; Go, S.J.; de Haan, A.; Wessels, J.G.H.  
Curr. Genet. 20, 137-144, 1991  
A:Title: Sequences of three dsRNAs associated with La France disease of the cultivated m  
A:Reference number: S17904; MUID:92035060; PMID:1934110  
A:Accession: S17905  
A>Status: preliminary  
A:Molecule type: mRNA

A;Residues: 1-359 <HAR>  
A;Cross-references: UNIPROT:Q83036; UNIPARC:UPI00000F4376; GB:D10829; GB:D00483; NID:922  
C;Superfamily: La France disease virus hypothetical protein M1

Query Match 24.3%; Score 49; DB 2; Length 359;  
Best Local Similarity 37.0%; Pred. No. 55;  
Matches 10; Conservative 5; Mismatches 10; Indels 2; Gaps 1;

Qy 7 CATQRLANFLVHSSNNFGPILPPTNVG 33  
Db 272 CKTELTEFTSGTNSSPGP--PPKRDG 296

## RESULT 48

B97852  
hypothetical protein RC1218 [imported] - Rickettsia conorii (strain Malish 7)  
C;Species: Rickettsia conorii  
C;Date: 30-Sep-2001 #sequence\_revision 30-Sep-2001 #text\_change 09-Jul-2004  
C;Accession: B97852  
R;Ogata, H.; Audic, S.; Reneato-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, D.; R  
Science 293, 2093-2098, 2001  
A;Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii.  
A;Reference number: A97700; MUID:21442074; PMID:11557893  
A;Accession: B97852  
A;Status: preliminary  
A;Molecule type: DNA  
A;Residues: 1-1162 <KUR>  
A;Cross-references: UNIPROT:Q92GA5; UNIPARC:UPI00000CC033; GB:AE006914; PIDN:AAL03756.1;  
C;Genetics:  
A;Gene: RC1218  
C;Superfamily: Rickettsia prowazekii hypothetical protein RP785

Query Match 24.3%; Score 49; DB 2; Length 1162;  
Best Local Similarity 35.5%; Pred. No. 1.8e+02;  
Matches 11; Conservative 2; Mismatches 18; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTN 31  
Db 840 KANTPICVLSDYTDRIEKPENIGPILVVD 870

## RESULT 49

S66525  
prostanoid receptor EP1 - mouse  
C;Species: Mus musculus (house mouse)  
C;Date: 28-Oct-1996 #sequence\_revision 13-Mar-1997 #text\_change 09-Jul-2004  
C;Accession: S66525; A48005  
R;Batschake, B.; Nilsson, C.; Sundelin, J.  
Eur. J. Biochem. 231, 809-814, 1995  
A;Title: Molecular characterization of the mouse prostanoid EP(1) receptor gene.  
A;Reference number: S66525; MUID:95377316; PMID:7649181  
A;Accession: S66525  
A;Molecule type: DNA  
A;Residues: 1-405 <BAT>  
A;Cross-references: UNIPROT:P35375; UNIPARC:UPI00000216D7; EMBL:Z49987; NID:G1197340; PI  
R;Watabe, A.; Sugimoto, Y.; Honda, A.; Irie, A.; Namba, T.; Negishi, M.; Ito, S.; Narumi  
J. Biol. Chem. 268, 20175-20178, 1993  
A;Title: Cloning and expression of cDNA for a mouse EP1 subtype of prostaglandin E recep  
A;Reference number: A48005; MUID:93388584; PMID:7690750  
A;Accession: A48005  
A;Molecule type: nucleic acid  
A;Residues: 1-405 <WAT>  
A;Cross-references: UNIPARC:UPI00000216D7; GB:D16338; NID:G439232; PIDN:BAA03842.1; PID:  
A;Experimental source: lung  
A;Note: sequence extracted from NCBI backbone (NCBIN:137716, NCBI:P:137717)  
C;Genetics:  
A;Introns: 317/3  
C;Superfamily: prostaglandin E receptor EP1  
C;Keywords: G protein-coupled receptor; glycoprotein; transmembrane protein  
F;40-62/Domain: transmembrane #status predicted <TM1>  
F;81-99/Domain: transmembrane #status predicted <TM2>  
F;114-135/Domain: transmembrane #status predicted <TM3>  
F;158-179/Domain: transmembrane #status predicted <TM4>

F;203-228/Domain: transmembrane #status predicted <TM5>  
F;302-323/Domain: transmembrane #status predicted <TM6>  
F;338-357/Domain: transmembrane #status predicted <TM7>

Query Match 24.0%; Score 48.5; DB 2; Length 405;  
Best Local Similarity 48.1%; Pred. No. 72;  
Matches 13; Conservative 0; Mismatches 11; Indels 3; Gaps 1;

Qy 5 ATCATQRLAN---FLVHSSNNFGPILP 28  
Db 15 ATCATPRLPNTSVVLPTGDNGTSPALP 41

## RESULT 50

T38730  
probable helicase - fission yeast (Schizosaccharomyces pombe)  
C;Species: Schizosaccharomyces pombe  
C;Date: 03-Dec-1999 #sequence\_revision 03-Dec-1999 #text\_change 09-Jul-2004  
C;Accession: T38730  
R;Gentles, S.; Churcher, C.M.; Barrell, B.G.; Rajandream, M.A.; Wood, V.  
submitted to the EMBL Data Library, September 1997  
A;Reference number: Z21797  
A;Accession: T38730  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-844 <GEN>  
A;Cross-references: UNIPROT:O14147; UNIPARC:UPI000006A4EE; EMBL:Z99167; PIDN:CAB16287.1;  
A;Experimental source: strain 972h-; cosmid c3G6  
C;Genetics:  
A;Gene: SPDB:SPAC3G6.11  
A;Map position: 1  
A;Introns: 213/3; 278/1

Query Match 24.0%; Score 48.5; DB 2; Length 844;  
Best Local Similarity 47.6%; Pred. No. 1.5e+02;  
Matches 10; Conservative 5; Mismatches 3; Indels 3; Gaps 1;

Qy 10 QRLANFL---VHSSNNFGPILP 27  
Db 810 RKLPNWLSKNHSSPNFGPAI 830

Search completed: May 12, 2006, 15:22:47  
Job time : 42 secs

GenCore version 5.1.1.8  
Copyright (c) 1993 - 2006 Bioceleration Ltd.

OM protein - protein search, using sw model

Run on: May 12, 2006, 15:22:21 ; Search time 48 Seconds  
(without alignments)  
63.729 Million cell updates/sec

Title: US-08-870-762B-1

Perfect score: 202

Sequence: 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 100 summaries

Database : Issued Patents AA:\*

1: /cgn2\_6/ptodata/1/1aa/5\_COMB.pep.\*

2: /cgn2\_6/ptodata/1/1aa/6\_COMB.pep.\*

3: /cgn2\_6/ptodata/1/1aa/H\_COMB.pep.\*

4: /cgn2\_6/ptodata/1/1aa/PCITUS\_COMB.pep.\*

5: /cgn2\_6/ptodata/1/1aa/RE\_COMB.pep.\*

6: /cgn2\_6/ptodata/1/1aa/backfiles1.pep.\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	202	100.0	37	1	US-08-477-727A-4
2	202	100.0	37	1	US-08-477-727A-29
3	202	100.0	37	1	US-08-477-727A-81
4	202	100.0	37	1	US-08-477-727A-87
5	202	100.0	37	1	US-08-471-675A-2
6	202	100.0	37	1	US-08-892-549-9
7	202	100.0	37	2	US-08-302-069A-1
8	202	100.0	37	2	US-09-576-062A-1
9	202	100.0	37	2	US-09-454-533-9
10	201	99.5	37	1	US-08-477-727A-84
11	201	99.5	37	1	US-08-477-727A-97
12	201	99.5	37	1	US-08-471-675A-8
13	201	99.5	37	1	US-08-471-675A-19
14	201	99.5	37	1	US-08-892-549-12
15	201	99.5	37	2	US-08-302-069A-7
16	201	99.5	37	2	US-09-576-062A-18
17	201	99.5	37	2	US-09-576-062A-7
18	201	99.5	37	2	US-09-576-062A-18
19	201	99.5	37	2	US-09-454-533-12
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21	199	98.5	37	1	US-08-477-727A-30
22	197	97.5	36	1	US-08-477-727A-88
23	197	97.5	36	1	US-08-471-675A-11
24	197	97.5	36	2	US-08-302-069A-10
25	197	97.5	36	2	US-09-576-062A-10
26	196	97.0	37	1	US-08-477-727A-77
27	195	96.5	37	1	US-08-477-727A-90

28	195	96.5	37	1	US-08-477-727A-96
29	195	96.5	37	1	US-08-471-675A-12
30	195	96.5	37	1	US-08-471-675A-18
31	195	96.5	37	1	US-08-892-549-16
32	195	96.5	37	1	US-08-892-549-22
33	195	96.5	37	2	US-08-302-069A-11
34	195	96.5	37	2	US-08-302-069A-17
35	195	96.5	37	2	US-09-576-062A-11
36	195	96.5	37	2	US-09-576-062A-17
37	195	96.5	37	2	US-09-454-533-16
38	195	96.5	37	2	US-09-454-533-22
39	194	96.0	37	1	US-08-471-675A-9
40	194	96.0	37	1	US-08-892-549-13
41	194	96.0	37	2	US-08-892-549-38
42	194	96.0	37	2	US-08-302-069A-8
43	194	96.0	37	2	US-09-576-062A-8
44	194	96.0	37	2	US-09-454-533-13
45	194	96.0	37	2	US-09-454-533-38
46	193	95.5	36	1	US-08-892-549-15
47	193	95.5	36	2	US-09-454-533-15
48	193	95.5	37	1	US-08-477-727A-17
49	193	95.5	37	1	US-08-477-727A-42
50	193	95.5	37	1	US-08-477-727A-89
51	192	95.0	36	1	US-08-892-549-40
52	192	95.0	36	2	US-09-454-533-40
53	192	95.0	37	1	US-08-477-727A-75
54	191	94.6	37	1	US-08-892-549-23
55	191	94.6	37	2	US-09-454-533-23
56	190	94.1	36	1	US-08-477-727A-98
57	190	94.1	36	1	US-08-471-675A-20
58	190	94.1	36	2	US-08-892-549-24
59	190	94.1	36	2	US-08-302-069A-19
60	190	94.1	36	2	US-09-576-062A-19
61	190	94.1	36	2	US-09-454-533-24
62	189	93.6	36	1	US-08-477-727A-86
63	189	93.6	36	1	US-08-892-549-39
64	189	93.6	36	2	US-08-302-069A-9
65	189	93.6	36	2	US-09-576-062A-9
66	189	93.6	36	2	US-09-454-533-39
67	188	93.1	37	1	US-08-477-727A-94
68	188	93.1	37	1	US-08-471-675A-16
69	188	93.1	37	1	US-08-892-549-20
70	188	93.1	37	2	US-08-302-069A-15
71	188	93.1	37	2	US-09-576-062A-15
72	188	93.1	37	2	US-09-454-533-20
73	187	92.6	37	1	US-07-794-288D-2
74	187	92.6	37	1	US-08-477-727A-91
75	187	92.6	37	1	US-08-471-675A-13
76	187	92.6	37	1	US-08-259-762-9
77	187	92.6	37	1	US-08-892-549-4
78	187	92.6	37	1	US-08-892-549-17
79	187	92.6	37	2	US-08-784-582-51
80	187	92.6	37	2	US-08-302-069A-12
81	187	92.6	37	2	US-09-070-504-17
82	187	92.6	37	2	US-09-576-062A-12
83	187	92.6	37	2	US-09-454-533-4
84	187	92.6	37	2	US-09-454-533-17
85	187	92.6	37	2	US-09-813-345C-17
86	187	92.6	37	2	US-09-623-548A-321
87	187	92.6	37	2	US-09-623-548A-334
88	187	92.6	37	2	US-09-657-276-321
89	187	92.6	37	2	US-09-657-276-334
90	187	92.6	93	2	US-08-589-028-8
91	187	92.6	93	2	US-08-784-582-8
92	187	92.6	93	2	US-08-785-271-8
93	186	92.1	37	1	US-08-477-727A-80
94	186	92.1	37	1	US-08-477-727A-82
95	186	92.1	37	1	US-08-477-727A-101
96	186	92.1	37	1	US-08-471-675A-4
97	186	92.1	37	1	US-08-471-675A-6
98	186	92.1	37	1	US-08-471-675A-23
99	186	92.1	37	1	US-08-892-549-8
100	186	92.1	37	1	US-08-892-549-10

ALIGNMENTS

```

RESULT 1
US-08-477-727A-4
; Sequence 4, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; US-08-477-727A-4

Query Match 100.0%; Score 202; DB 1; Length 37;
Best Local Similarity 100.0%; Pred. No. 6e-21;
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 2
US-08-477-727A-29
; Sequence 29, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
```

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; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 29:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; US-08-477-727A-29

Query Match 100.0%; Score 202; DB 1; Length 37;
Best Local Similarity 100.0%; Pred. No. 6e-21;
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 3
US-08-477-727A-81
; Sequence 81, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
```



COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSEQ Version 1.5  
CURRENT APPLICATION DATA: US/08/477,727A  
FILING DATE: 07-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 214/005  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 81:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
FRAGMENT TYPE: internal  
ORIGINAL SOURCE:  
US-08-477-727A-81

Query Match 100.0%; Score 202; DB 1; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 4  
US-08-477-727A-87  
Sequence 87, Application US/08477727A  
Patent No. 5739106  
GENERAL INFORMATION:  
APPLICANT: Rink, Timothy  
APPLICANT: Young, Andrew  
APPLICANT: Beeley, Nigel  
APPLICANT: Prickett, Kathryn  
TITLE OF INVENTION: APPETITE REGULATING  
COMPOSITIONS  
NUMBER OF SEQUENCES: 108  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSEQ Version 1.5  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/477,727A  
FILING DATE: 07-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:

ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 214/005  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 87:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
FRAGMENT TYPE: internal  
ORIGINAL SOURCE:  
US-08-477-727A-87

Query Match 100.0%; Score 202; DB 1; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 5  
US-08-471-675A-2  
Sequence 2, Application US/08471675A  
Patent No. 5795861  
GENERAL INFORMATION:  
APPLICANT: Kolterman, Orville  
APPLICANT: Rink, Timothy  
TITLE OF INVENTION: METHODS FOR REGULATING  
GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSEQ Version 2.0  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/471,675A  
FILING DATE: 05-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/302,069  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 213/048  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid

STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
OTHER INFORMATION: disulfide bridge between the Cys  
OTHER INFORMATION: residues at positions 2 and 7;  
OTHER INFORMATION: amidated Tyr at position 37  
US-08-471-675A-2

Query Match 100.0%; Score 202; DB 1; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 6

US-08-892-549-9  
Sequence 9, Application US/08892549  
Patent No. 5998367  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
TITLE OF INVENTION: USES THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/892,549  
FILING DATE: 14-JUL-1997  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/447,849  
FILING DATE: 23-MAY-1995  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 9:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)

US-08-892-549-9

Query Match 100.0%; Score 202; DB 1; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 7

US-08-302-069A-1  
Sequence 1, Application US/08302069A  
Patent No. 6114304  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: RINK, Timothy J.

APPLICANT: BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
CLASSIFICATION: 514

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)

Query Match 100.0%; Score 202; DB 2; Length 37;  
Best Local Similarity 100.0%; Pred. No. 6e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

## RESULT 8

```
US-09-576-062A-1
; Sequence 1, Application US/09576062A
; Patent No. 6608029
; GENERAL INFORMATION:
; APPLICANT: KOLTERMAN, Orville G.
; RINK, Timothy J.
; BROWN, Kathleen Ann Keiting
; TITLE OF INVENTION: METHODS FOR REGULATING
; GASTROINTESTINAL MOTILITY
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/576,062A
; FILING DATE: 22-May-2000
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/302,069
; FILING DATE: 07-SEP-1994
; APPLICATION NUMBER: 08/118,381
; FILING DATE: 07-SEP-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 209/146
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-576-062A-1
Query Match 100.0%; Score 202; DB 2; Length 37;
Best Local Similarity 100.0%; Pred. No. 6e-21;
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
RESULT 9
US-09-454-533-9
; Sequence 9, Application US/09454533
; Patent No. 6610824
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
US-08-870-727A-84
; Sequence 84, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
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; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ Version 1.5
; CURRENT APPLICATION DATA: US/08/477.727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 84:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; US-08-477-727A-84

Query Match          99.5%; Score 201; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 8.3e-21;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 11
US-08-477-727A-97
; Sequence 97, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477.727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 84:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; US-08-477-727A-97

Query Match          99.5%; Score 201; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 8.3e-21;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 12
US-08-471-675A-8
; Sequence 8, Application US/08471675A
; Patent No. 5795861
; GENERAL INFORMATION:
; APPLICANT: Kolterman, Orville
; APPLICANT: Rink, Timothy
; TITLE OF INVENTION: METHODS FOR REGULATING
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/471.675A
; FILING DATE: 05-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 213/048
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
```

FEATURE: disulfide bridge between the Cys  
OTHER INFORMATION: residues at positions 2 and 7;  
OTHER INFORMATION: amidated Tyr at position 37  
US-08-471-675A-8

Query Match 99.5%; Score 201; DB 1; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||

Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||

RESULT 13

US-08-471-675A-19  
Sequence 19, Application US/08471675A  
Patent No. 5795861  
GENERAL INFORMATION:  
APPLICANT: Kolterman, Orville  
APPLICANT: Rink, Timothy  
TITLE OF INVENTION: METHODS FOR REGULATING  
TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066

COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSeq version 2.0  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/471,675A  
FILING DATE: 05-JUN-1995  
CLASSIFICATION: 514

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/302,069  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 213/048  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:

INFORMATION FOR SEQ ID NO: 19:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:

OTHER INFORMATION: disulfide bridge between the Cys  
OTHER INFORMATION: residues at positions 2 and 7;  
OTHER INFORMATION: amidated Tyr at position 37  
US-08-471-675A-19

Query Match 99.5%; Score 201; DB 1; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||

Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 14

US-08-892-549-12  
Sequence 12, Application US/08892549  
Patent No. 5998367  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
TITLE OF INVENTION: USES THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/892,549  
FILING DATE: 14-JUL-1997  
CLASSIFICATION: 514

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/447,849  
FILING DATE: 23-MAY-1995  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 12:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-892-549-12

Query Match 99.5%; Score 201; DB 1; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||

Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||

RESULT 15

US-08-302-069A-7  
Sequence 7, Application US/08302069A  
Patent No. 6114304  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.

APPLICANT: YOUNG, Andrew A.  
APPLICANT: RINK, Timothy J.  
APPLICANT: BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 7:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-302-069A-7

Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 16  
US-08-302-069A-18  
Sequence 18, Application US/08302069A  
Patent No. 6114304  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: RINK, Timothy J.  
APPLICANT: BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES

STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 18:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 2,7  
OTHER INFORMATION: disulfide bridge between  
OTHER INFORMATION: the Cys residues  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-302-069A-18

Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 17  
US-09-576-062A-7  
Sequence 7, Application US/09576062A  
Patent No. 6608029  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: RINK, Timothy J.  
APPLICANT: BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/576,062A

;/ FILING DATE: 22-May-2000  
;/ CLASSIFICATION: <Unknown>  
;/ PRIOR APPLICATION DATA:  
;/ APPLICATION NUMBER: 08/302,069  
;/ FILING DATE: 07-SEP-1994  
;/ APPLICATION NUMBER: 08/118,381  
;/ FILING DATE: 07-SEP-1993  
;/ ATTORNEY/AGENT INFORMATION:  
;/ NAME: DUFT, BRADFORD J.  
;/ REGISTRATION NUMBER: 32,219  
;/ REFERENCE/DOCKET NUMBER: 209/146  
;/ TELECOMMUNICATION INFORMATION:  
;/ TELEPHONE: 619/552-2200  
;/ TELEFAX: 213/955-0440  
;/ TELEX: 67-3510  
;/ INFORMATION FOR SEQ ID NO: 7:  
;/ SEQUENCE CHARACTERISTICS:  
;/ LENGTH: 37 amino acids  
;/ TYPE: amino acid  
;/ STRANDEDNESS: single  
;/ TOPOLOGY: linear  
;/ MOLECULE TYPE: peptide  
;/ FEATURE:  
;/ LOCATION: 37  
;/ OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
;/ SEQUENCE DESCRIPTION: SEQ ID NO: 7:  
US-09-576-062A-7  
  
Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
  
RESULT 18  
US-09-576-062A-18  
;/ Sequence 18, Application US/09576062A  
;/ Patent No. 6608029  
;/ GENERAL INFORMATION:  
;/ APPLICANT: KOLTERMAN, Orville G.  
;/ YOUNG, Andrew A.  
;/ RINK, Timothy J.  
;/ BROWN, Kathleen Ann Keiting  
;/ TITLE OF INVENTION: METHODS FOR REGULATING  
;/ NUMBER OF SEQUENCES: 30  
;/ CORRESPONDENCE ADDRESS:  
;/ ADDRESSEE: LYON & LYON  
;/ STREET: 633 WEST FIFTH STREET  
;/ CITY: LOS ANGELES  
;/ STATE: CALIFORNIA  
;/ COUNTRY: USA  
;/ ZIP: 90017  
;/ COMPUTER READABLE FORM:  
;/ MEDIUM TYPE: Floppy disk  
;/ COMPUTER: IBM PC compatible  
;/ OPERATING SYSTEM: PC-DOS/MS-DOS  
;/ SOFTWARE: Patentin Release #1.0, Version #1.25  
;/ CURRENT APPLICATION NUMBER: US/09/576,062A  
;/ FILING DATE: 22-May-2000  
;/ CLASSIFICATION: <Unknown>  
;/ PRIOR APPLICATION DATA:  
;/ APPLICATION NUMBER: 08/302,069  
;/ FILING DATE: 07-SEP-1994  
;/ APPLICATION NUMBER: 08/118,381  
;/ FILING DATE: 07-SEP-1993  
;/ ATTORNEY/AGENT INFORMATION:  
;/ NAME: DUFT, BRADFORD J.  
;/ REGISTRATION NUMBER: 32,219

;/ REFERENCE/DOCKET NUMBER: 209/146  
;/ TELECOMMUNICATION INFORMATION:  
;/ TELEPHONE: 619/552-2200  
;/ TELEFAX: 213/955-0440  
;/ TELEX: 67-3510  
;/ INFORMATION FOR SEQ ID NO: 18:  
;/ SEQUENCE CHARACTERISTICS:  
;/ LENGTH: 37 amino acids  
;/ TYPE: amino acid  
;/ STRANDEDNESS: single  
;/ TOPOLOGY: linear  
;/ MOLECULE TYPE: protein  
;/ FEATURE:  
;/ LOCATION: 37  
;/ OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
;/ SEQUENCE DESCRIPTION: SEQ ID NO: 18:  
US-09-576-062A-18  
  
Query Match 99.5%; Score 201; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 8.3e-21;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
  
RESULT 19  
US-09-454-533-12  
;/ Sequence 12, Application US/09454533  
;/ Patent No. 6610824  
;/ GENERAL INFORMATION:  
;/ APPLICANT: GAETA, Laura S.L. Et Al.  
;/ TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
;/ NUMBER OF SEQUENCES: 41  
;/ CORRESPONDENCE ADDRESS:  
;/ ADDRESSEE: LYON & LYON  
;/ STREET: 633 WEST FIFTH STREET  
;/ CITY: LOS ANGELES  
;/ STATE: CALIFORNIA  
;/ COUNTRY: USA  
;/ ZIP: 90017  
;/ COMPUTER READABLE FORM:  
;/ MEDIUM TYPE: Floppy disk  
;/ COMPUTER: IBM PC compatible  
;/ OPERATING SYSTEM: PC-DOS/MS-DOS  
;/ SOFTWARE: Patentin Release #1.0, Version #1.25  
;/ CURRENT APPLICATION NUMBER: US/09/454,533  
;/ FILING DATE: 06-Dec-1999  
;/ CLASSIFICATION: <Unknown>  
;/ PRIOR APPLICATION DATA:  
;/ APPLICATION NUMBER: 08/892,549  
;/ FILING DATE: <Unknown>  
;/ APPLICATION NUMBER: 07/794,266  
;/ FILING DATE: 19-NOV-1991  
;/ APPLICATION NUMBER: US 07/667,040  
;/ FILING DATE: 08-MAR-1991  
;/ ATTORNEY/AGENT INFORMATION:  
;/ NAME: DUFT, BRADFORD J.  
;/ REGISTRATION NUMBER: 32,219  
;/ REFERENCE/DOCKET NUMBER: 227/006  
;/ TELECOMMUNICATION INFORMATION:  
;/ TELEPHONE: 619/552-2200  
;/ TELEFAX: 213/955-0440  
;/ TELEX: 67-3510  
;/ INFORMATION FOR SEQ ID NO: 12:  
;/ SEQUENCE CHARACTERISTICS:  
;/ LENGTH: 37 amino acids  
;/ TYPE: amino acid  
;/ STRANDEDNESS: single  
;/ TOPOLOGY: linear

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; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-454-533-12
Query Match      99.5%; Score 201; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 8.3e-21;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 20
US-08-477-727A-5
; Sequence 5, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-08-477-727A-5
Query Match      98.5%; Score 199; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.6e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 22
US-08-477-727A-88
; Sequence 88, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-08-477-727A-88
Query Match      98.5%; Score 199; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.6e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 21
US-08-477-727A-30
; Sequence 30, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 30:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-08-477-727A-30
Query Match      98.5%; Score 199; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.6e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 22
US-08-477-727A-88
; Sequence 88, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; TITLE OF INVENTION: COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 30:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-08-477-727A-88
Query Match      98.5%; Score 199; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 1.6e-20;
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
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APPLICANT: Prickett, Kathryn  
TITLE OF INVENTION: APPETITE REGULATING  
COMPOSITIONS  
NUMBER OF SEQUENCES: 108  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSEQ Version 1.5  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/477,727A  
FILING DATE: 07-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 214/005  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 88:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
FRAGMENT TYPE: internal  
ORIGINAL SOURCE:  
US-08-477-727A-88

Query Match 97.5%; Score 197; DB 1; Length 36;  
Best Local Similarity 100.0%; Pred. No. 2.9e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 23  
US-08-471-675A-11  
Sequence 11, Application US/08471675A  
Patent No. 5795861  
GENERAL INFORMATION:  
APPLICANT: Koltzman, Orville  
APPLICANT: Rink, Timothy  
TITLE OF INVENTION: METHODS FOR REGULATING  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSES: LYON & LYON  
STREET: 633 WEST FIFTH STREET, SUITE 4700  
CITY: LOS ANGELES  
STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS  
SOFTWARE: FastSEQ Version 2.0  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/471,675A  
FILING DATE: 05-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/302,069  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 213/048  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
OTHER INFORMATION: disulfide bridge between the Cys  
residues at positions 1 and 6;  
OTHER INFORMATION: amidated Tyr at position 36  
US-08-471-675A-11

Query Match 97.5%; Score 197; DB 1; Length 36;  
Best Local Similarity 100.0%; Pred. No. 2.9e-20;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 24  
US-08-302-069A-10  
Sequence 10, Application US/08302069A  
Patent No. 6114304  
GENERAL INFORMATION:  
APPLICANT: KOLTERMAN, Orville G.  
APPLICANT: YOUNG, Andrew A.  
APPLICANT: RINK, Timothy J.  
APPLICANT: BROWN, Kathleen Ann Keiting  
TITLE OF INVENTION: METHODS FOR REGULATING  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/302,069A  
FILING DATE: 07-SEP-1994  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:

```
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 209/146
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 1,6
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 36
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; US-08-302-069A-10

Query Match 97.5%; Score 197; DB 2; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.9e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 25
US-08-576-062A-10
; Sequence 10, Application US/09576062A
; Patent No. 6608029
; GENERAL INFORMATION:
; APPLICANT: KOLTERMAN, Orville G.
; YOUNG, Andrew A.
; RINK, Timothy J.
; BROWN, Kathleen Ann Keiting
; TITLE OF INVENTION: METHODS FOR REGULATING
; GASTROINTESTINAL MOTILITY
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/576,062A
; FILING DATE: 22-May-2000
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/302,069
; FILING DATE: 07-SEP-1994
; APPLICATION NUMBER: 08/118,381
; FILING DATE: 07-SEP-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 209/146
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 10:
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; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 36
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; US-09-576-062A-10

Query Match 97.5%; Score 197; DB 2; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.9e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 CNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 36

RESULT 26
US-08-477-727A-77
; Sequence 77, Application US/08477727A
; Patent No. 5739106
; GENERAL INFORMATION:
; APPLICANT: Rink, Timothy
; APPLICANT: Young, Andrew
; APPLICANT: Beeley, Nigel
; APPLICANT: Prickett, Kathryn
; TITLE OF INVENTION: APPETITE REGULATING
; COMPOSITIONS
; NUMBER OF SEQUENCES: 108
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 1.5
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/477,727A
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 214/005
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 77:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; MOLECULE TYPE: peptide
; TOPOLOGY: linear
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; US-08-477-727A-77
```

Query Match 97.0%; Score 196; DB 1; Length 37;  
Best Local Similarity 97.3%; Pred. No. 4.1e-20;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNDY 37

## RESULT 27

US-08-477-727A-90  
; Sequence 90, Application US/08477727A  
; Patent No. 5739106  
; GENERAL INFORMATION:  
; APPLICANT: Rink, Timothy  
; APPLICANT: Young, Andrew  
; APPLICANT: Beeley, Nigel  
; APPLICANT: Prickett, Kathryn  
; TITLE OF INVENTION: APPETITE REGULATING  
; TITLE OF INVENTION: COMPOSITIONS  
; NUMBER OF SEQUENCES: 108  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 90071-2066  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSEQ Version 1.5  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/477,727A  
; FILING DATE: 07-JUN-1995  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER:  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 214/005  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619-552-8400  
; TELEFAX: 619-552-0157  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 90:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; HYPOTHETICAL: NO  
; ANTI-SENSE: NO  
; FRAGMENT TYPE: internal  
; ORIGINAL SOURCE:  
; US-08-477-727A-90

Query Match 96.5%; Score 195; DB 1; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

## RESULT 28

US-08-477-727A-96  
; Sequence 96, Application US/08477727A

; Patent No. 5739106  
; GENERAL INFORMATION:  
; APPLICANT: Rink, Timothy  
; APPLICANT: Young, Andrew  
; APPLICANT: Beeley, Nigel  
; APPLICANT: Prickett, Kathryn  
; TITLE OF INVENTION: APPETITE REGULATING  
; TITLE OF INVENTION: COMPOSITIONS  
; NUMBER OF SEQUENCES: 108  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 90071-2066  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSEQ Version 1.5  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/477,727A  
; FILING DATE: 07-JUN-1995  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER:  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 214/005  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619-552-8400  
; TELEFAX: 619-552-0157  
; TELEX:  
; INFORMATION FOR SEQ ID NO: 96:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; HYPOTHETICAL: NO  
; ANTI-SENSE: NO  
; FRAGMENT TYPE: internal  
; ORIGINAL SOURCE:  
; US-08-477-727A-96

Query Match 96.5%; Score 195; DB 1; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

## RESULT 29

US-08-471-675A-12  
; Sequence 12, Application US/08471675A  
; Patent No. 5795861  
; GENERAL INFORMATION:  
; APPLICANT: Kolterman, Orville  
; APPLICANT: Rink, Timothy  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA

;; COUNTRY: USA  
;; ZIP: 90071-2066  
;; COMPUTER READABLE FORM:  
;; MEDIUM TYPE: Diskette  
;; COMPUTER: IBM Compatible  
;; OPERATING SYSTEM: DOS  
;; SOFTWARE: FastSEQ Version 2.0  
;; CURRENT APPLICATION DATA:  
;; APPLICATION NUMBER: US/08/471.675A  
;; FILING DATE: 05-JUN-1995  
;; CLASSIFICATION: 514  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: 08/302,069  
;; FILING DATE: 07-SEP-1994  
;; APPLICATION NUMBER: 08/118,381  
;; FILING DATE: 07-SEP-1993  
;; NAME: DUFT, BRADFORD J  
;; ATTORNEY/AGENT INFORMATION:  
;; REGISTRATION NUMBER: 32,219  
;; REFERENCE/DOCKET NUMBER: 213/048  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: 619-552-8400  
;; TELEFAX: 619-552-0157  
;; TELEX:  
;; INFORMATION FOR SEQ ID NO: 12:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 37 amino acids  
;; TYPE: amino acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: peptide  
;; FEATURE:  
;; OTHER INFORMATION: disulfide bridge between the Cys  
;; OTHER INFORMATION: residues at positions 2 and 7;  
;; OTHER INFORMATION: amidated Tyr at position 37  
US-08-471-675A-12

Query Match 96.5%; Score 195; DB 1; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37  
|||||

RESULT 30  
US-08-471-675A-18  
; Sequence 18, Application US/08471675A  
; Patent No. 5795861  
; GENERAL INFORMATION:  
; APPLICANT: Kolterman, Orville  
; APPLICANT: Rink, Timothy  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; TITLE OF INVENTION: GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET, SUITE 4700  
; CITY: LOS ANGELES  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 90071-2066  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSEQ Version 2.0  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/471.675A  
; FILING DATE: 05-JUN-1995  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:

;; APPLICATION NUMBER: 08/302,069  
;; FILING DATE: 07-SEP-1994  
;; APPLICATION NUMBER: 08/118,381  
;; FILING DATE: 07-SEP-1993  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: DUFT, BRADFORD J  
;; REGISTRATION NUMBER: 32,219  
;; REFERENCE/DOCKET NUMBER: 213/048  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: 619-552-8400  
;; TELEFAX: 619-552-0157  
;; TELEX:  
;; INFORMATION FOR SEQ ID NO: 18:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 37 amino acids  
;; TYPE: amino acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: peptide  
;; FEATURE:  
;; OTHER INFORMATION: disulfide bridge between the Cys  
;; OTHER INFORMATION: residues at positions 2 and 7;  
;; OTHER INFORMATION: amidated Tyr at position 37  
US-08-471-675A-18

Query Match 96.5%; Score 195; DB 1; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37  
|||||

RESULT 31  
US-08-892-549-16  
; Sequence 16, Application US/08892549  
; Patent No. 5998367  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et Al.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; TITLE OF INVENTION: USES THEREFOR  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/892,549  
; FILING DATE: 14-JUL-1997  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/447,849  
; FILING DATE: 23-MAY-1995  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-NOV-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-MAR-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 227/006  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440

```

;
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 16:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
;
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
;
US-08-892-549-16

Query Match 96.5%; Score 195; DB 1; Length 37;
Best Local Similarity 94.6%; Pred. No. 5.6e-20;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCVTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
Db 1 KCVTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 32
US-08-892-549-22
; Sequence 22, Application US/08892549
; Patent No. 5998367
; GENERAL INFORMATION:
; APPLICANT: GASTA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; TITLE OF INVENTION: USES THEREOF
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/892,549
; FILING DATE: 14-JUL-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/447,849
; FILING DATE: 23-MAY-1995
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 22:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
```

```

;
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
;
US-08-892-549-22

Query Match 96.5%; Score 195; DB 1; Length 37;
Best Local Similarity 94.6%; Pred. No. 5.6e-20;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KCVTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
Db 1 KCVTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 33
US-08-302-069A-11
; Sequence 11, Application US/08302069A
; Patent No. 6114304
; GENERAL INFORMATION:
; APPLICANT: KOLTERMAN, Orville G.
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: BROWN, Kathleen Ann Keiting
; TITLE OF INVENTION: METHODS FOR REGULATING
; TITLE OF INVENTION: GASTROINTESTINAL MOTILITY
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/302,069A
; FILING DATE: 07-SEP-1994
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/118,381
; FILING DATE: 07-SEP-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 209/146
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
;
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
;
US-08-302-069A-11

Query Match 96.5%; Score 195; DB 2; Length 37;
Best Local Similarity 94.6%; Pred. No. 5.6e-20;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
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Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37

## RESULT 34

US-08-302-069A-17  
; Sequence 17, Application US/08302069A  
; Patent No. 6114304  
; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; APPLICANT: YOUNG, Andrew A.  
; APPLICANT: RINK, Timothy J.  
; APPLICANT: BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/302.069A  
; FILING DATE: 07-SEP-1994  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/118,381  
; FILING DATE: 07-SEP-1993  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 209/146  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 17:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; FEATURE:  
; LOCATION: 2,7  
; OTHER INFORMATION: disulfide bridge between  
; OTHER INFORMATION: the Cys residues  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-302-069A-17

Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37

## RESULT 35

US-09-576-062A-11  
; Sequence 11, Application US/09576062A  
; Patent No. 6608029

; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; YOUNG, Andrew A.  
; RINK, Timothy J.  
; BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/576.062A  
; FILING DATE: 22-May-2000  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/302,069  
; FILING DATE: 07-SEP-1994  
; APPLICATION NUMBER: 08/118,381  
; FILING DATE: 07-SEP-1993  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 209/146  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 11:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 37 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FEATURE:  
; LOCATION: 37  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 11:  
US-09-576-062A-11  
  
Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
Qy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
Db 1 KCNTATCATQRLANFLVHSSNNLGPVLPPTNVGSNTY 37  
  
RESULT 36  
US-09-576-062A-17  
; Sequence 17, Application US/09576062A  
; Patent No. 6608029  
; GENERAL INFORMATION:  
; APPLICANT: KOLTERMAN, Orville G.  
; YOUNG, Andrew A.  
; RINK, Timothy J.  
; BROWN, Kathleen Ann Keiting  
; TITLE OF INVENTION: METHODS FOR REGULATING  
; GASTROINTESTINAL MOTILITY  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON

STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/576,062A  
FILING DATE: 22-May-2000  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/302,069  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 17:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 17:  
US-09-576-062A-17

Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37  
|||||  
RESULT 37  
US-09-454-533-16  
Sequence 16, Application US/09454533  
Patent No. 6610824  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USSES THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999

CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 16:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 16:  
US-09-454-533-16  
Query Match 96.5%; Score 195; DB 2; Length 37;  
Best Local Similarity 94.6%; Pred. No. 5.6e-20;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNNLGPILPPTNVGSNTY 37  
|||||  
RESULT 38  
US-09-454-533-22  
Sequence 22, Application US/09454533  
Patent No. 6610824  
GENERAL INFORMATION:  
APPLICANT: GAETA, Laura S.L. Et Al.  
TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
USSES THEREFOR  
NUMBER OF SEQUENCES: 41  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/454,533  
FILING DATE: 06-Dec-1999  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/892,549  
FILING DATE: <Unknown>  
APPLICATION NUMBER: 07/794,266  
FILING DATE: 19-NOV-1991  
APPLICATION NUMBER: US 07/667,040  
FILING DATE: 08-MAR-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219

```
;
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 22:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
; SEQUENCE DESCRIPTION: SEQ ID NO: 22:
US-09-454-533-22
Query Match 96.5%; Score 195; DB 2; Length 37;
Best Local Similarity 94.6%; Pred. No. 5.6e-20;
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 39
US-08-471-675A-9
; Sequence 9, Application US/08471675A
; Patent No. 5795861
; GENERAL INFORMATION:
; APPLICANT: Kolterman, Orville
; APPLICANT: Rink, Timothy
; TITLE OF INVENTION: METHODS FOR REGULATING
; TITLE OF INVENTION: GASTROINTESTINAL MOTILITY
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET, SUITE 4700
; CITY: LOS ANGELES
; STATE: CA
; COUNTRY: USA
; ZIP: 90071-2066
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/471,675A
; FILING DATE: 05-JUN-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/302,069
; FILING DATE: 07-SEP-1994
; APPLICATION NUMBER: 08/118,381
; FILING DATE: 07-SEP-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 213/048
; TELEPHONE: 619-552-8400
; TELEFAX: 619-552-0157
; TELEX:
; INFORMATION FOR SEQ ID NO: 9:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
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; FEATURE:
; OTHER INFORMATION: disulfide bridge between the Cys
; OTHER INFORMATION: residues at positions 2 and 7;
; OTHER INFORMATION: amidated Tyr at position 37
US-08-471-675A-9
Query Match 96.0%; Score 194; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 7.7e-20;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 1 KCNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37
| | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 40
US-08-892-549-13
; Sequence 13, Application US/08892549
; Patent No. 5998367
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; TITLE OF INVENTION: USES THEREFOR
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/892,549
; FILING DATE: 14-JUL-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/447,849
; FILING DATE: 23-MAY-1995
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
US-08-892-549-13
Query Match 96.0%; Score 194; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 7.7e-20;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 41
US-08-892-549-38
; Sequence 38, Application US/08892549
; Patent No. 5998367
; GENERAL INFORMATION:
; APPLICANT: GAETA, Laura S.L. Et Al.
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/892,549
; FILING DATE: 14-JUL-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/447,849
; FILING DATE: 23-MAY-1995
; APPLICATION NUMBER: 07/794,266
; FILING DATE: 19-NOV-1991
; APPLICATION NUMBER: US 07/667,040
; FILING DATE: 08-MAR-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 227/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 38:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
US-08-892-549-38

Query Match 96.0%; Score 194; DB 1; Length 37;
Best Local Similarity 97.3%; Pred. No. 7.7e-20;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 42
US-08-302-069A-8
; Sequence 8, Application US/08302069A
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```
; Patent No. 6114304
; GENERAL INFORMATION:
; APPLICANT: KOLTERMAN, Orville G.
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: RINK, Timothy J.
; APPLICANT: BROWN, Kathleen Ann Keiting
; TITLE OF INVENTION: METHODS FOR REGULATING
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSES: LYON & LYON
; STREET: 633 WEST FIFTH STREET
; CITY: LOS ANGELES
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 90017
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/302,069A
; FILING DATE: 07-SEP-1994
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/118,381
; FILING DATE: 07-SEP-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: DUFT, BRADFORD J.
; REGISTRATION NUMBER: 32,219
; REFERENCE/DOCKET NUMBER: 209/146
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 619/552-2200
; TELEFAX: 213/955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 37 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FEATURE:
; LOCATION: 2,7
; OTHER INFORMATION: disulfide bridge between
; OTHER INFORMATION: the Cys residues
; LOCATION: 37
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)
US-08-302-069A-8

Query Match 96.0%; Score 194; DB 2; Length 37;
Best Local Similarity 97.3%; Pred. No. 7.7e-20;
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37
Db 1 KNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 43
US-09-576-062A-8
; Sequence 8, Application US/09576062A
; Patent No. 6608029
; GENERAL INFORMATION:
; APPLICANT: KOLTERMAN, Orville G.
; APPLICANT: YOUNG, Andrew A.
; APPLICANT: RINK, Timothy J.
; APPLICANT: BROWN, Kathleen Ann Keiting
; TITLE OF INVENTION: METHODS FOR REGULATING
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
```

ADDRESSEE: LYON & LYON  
STREET: 633 WEST FIFTH STREET  
CITY: LOS ANGELES  
STATE: CALIFORNIA  
COUNTRY: USA  
ZIP: 90017  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/576,062A  
FILING DATE: 22-MAY-2000  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/302,069  
FILING DATE: 07-SEP-1994  
APPLICATION NUMBER: 08/118,381  
FILING DATE: 07-SEP-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J.  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 209/146  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 8:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 8:  
US-09-576-062A-8

Query Match 96.0%; Score 194; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 7.7e-20;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 44  
US-09-454-533-13  
; Sequence 13, Application US/09454533  
; Patent No. 6610824  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et Al.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; USES THEREFOR  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/454,533

; FILING DATE: 06-Dec-1999  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/892,549  
; FILING DATE: <Unknown>  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-NOV-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-MAR-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 227/006  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 13:  
; SEQUENCE DESCRIPTION: SEQ ID NO: 13:  
US-09-454-533-13

Query Match 96.0%; Score 194; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 7.7e-20;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
DB 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37

RESULT 45  
US-09-454-533-38  
; Sequence 38, Application US/09454533  
; Patent No. 6610824  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et Al.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; USES THEREFOR  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; STREET: 633 WEST FIFTH STREET  
; CITY: LOS ANGELES  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/454,533  
; FILING DATE: 06-Dec-1999  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/892,549  
; FILING DATE: <Unknown>  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-NOV-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-MAR-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.

REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 227/006  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619/552-2200  
TELEFAX: 213/955-0440  
TELEX: 67-3510  
INFORMATION FOR SEQ ID NO: 38  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 37  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
SEQUENCE DESCRIPTION: SEQ ID NO: 38:  
US-09-454-533-38

Query Match 96.0%; Score 194; DB 2; Length 37;  
Best Local Similarity 97.3%; Pred. No. 7.7e-20;  
Matches 36; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 KNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37

RESULT 46  
US-08-892-549-15  
; Sequence 15, Application US/08892549  
; Patent No. 5998367  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et AL.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; CITY: 633 WEST FIFTH STREET  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/892,549  
; FILING DATE: 14-JUL-1997  
; CLASSIFICATION: 514  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/447,849  
; FILING DATE: 23-MAY-1995  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-NOV-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-MAR-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 227/006  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 15:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 36 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single

TOPOLOGY: linear  
MOLECULE TYPE: protein  
FEATURE:  
LOCATION: 1,6  
OTHER INFORMATION: disulfide bridge between  
the Cys residues  
LOCATION: 36  
OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
US-08-892-549-15  
Query Match 95.5%; Score 193; DB 1; Length 36;  
Best Local Similarity 97.2%; Pred. No. 1e-19;  
Matches 35; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 37  
DB 1 CNTATCATQRLANFLVHSSNFGPILPPTNVGSNTY 36

RESULT 47  
US-09-454-533-15  
; Sequence 15, Application US/09454533  
; Patent No. 6610824  
; GENERAL INFORMATION:  
; APPLICANT: GAETA, Laura S.L. Et AL.  
; TITLE OF INVENTION: NOVEL AMYLIN AGONIST PEPTIDES AND  
; NUMBER OF SEQUENCES: 41  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LYON & LYON  
; CITY: 633 WEST FIFTH STREET  
; STATE: CALIFORNIA  
; COUNTRY: USA  
; ZIP: 90017  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/454,533  
; FILING DATE: 06-Dec-1999  
; CLASSIFICATION: <Unknown>  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: 08/892,549  
; FILING DATE: <Unknown>  
; APPLICATION NUMBER: 07/794,266  
; FILING DATE: 19-NOV-1991  
; APPLICATION NUMBER: US 07/667,040  
; FILING DATE: 08-MAR-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: DUFT, BRADFORD J.  
; REGISTRATION NUMBER: 32,219  
; REFERENCE/DOCKET NUMBER: 227/006  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 619/552-2200  
; TELEFAX: 213/955-0440  
; TELEX: 67-3510  
; INFORMATION FOR SEQ ID NO: 15:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 36 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; FEATURE:  
; LOCATION: 36  
; OTHER INFORMATION: amidated Tyr (Tyrosinamide)  
; SEQUENCE DESCRIPTION: SEQ ID NO: 15:  
US-09-454-533-15  
Query Match 95.5%; Score 193; DB 2; Length 36;



STATE: CA  
COUNTRY: USA  
ZIP: 90071-2066  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: DOS  
SOFTWARE: FastSeq Version 1.5  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/477,727A  
FILING DATE: 07-JUN-1995  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: DUFT, BRADFORD J  
REGISTRATION NUMBER: 32,219  
REFERENCE/DOCKET NUMBER: 214/005  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-8400  
TELEFAX: 619-552-0157  
TELEX:  
INFORMATION FOR SEQ ID NO: 89:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 37 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
FRAGMENT TYPE: internal  
ORIGINAL SOURCE:  
US-08-477-727A-89

Query Match 95.5%; Score 193; DB 1; Length 37;  
Best Local Similarity 94.6%; Pred. No. 1.1e-19;  
Matches 35; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Oy 1 KCNTATCATQRLANFLVHSSNNFGPILPPTNVGSNTY 37  
|||||  
Db 1 KCNTATCATQRLANFLVHSSNNFGPVLPTNVGSNTY 37  
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Search completed: May 12, 2006, 15:23:38  
Job time : 49 secs

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